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Sir:

Enclosed herewith for filing is a patent application of Wilson, et al. entitled SCREENING METHODS FOR COMPOUNDS USEFUL IN THE TREATMENT OF POLYCYSTIC KIDNEY DISEASE.

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☐ Priority of application Country __, Appln. No. __ filed __ is claimed under 35 U.S.C. 119.

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TO ALL WHOM IT MAY CONCERN:

Be it known that WE, PATRICIA WILSON and CHRIS BURROW, citizens of the United Kingdom and the United States of America, respectively, whose post office addresses are 525 East 80th Street, Apt #8D, New York, N.Y. 10021 have invented an improvement in

SCREENING METHODS FOR COMPOUNDS USEFUL IN THE
TREATMENT OF POLYCYSTIC KIDNEY DISEASE

of which the following is a

SPECIFICATION

1. INTRODUCTION

The present invention provides cell-based screening assays designed to identify agents that regulate the activity of the polycystic kidney disease proteins encoded by the PKD-1 and PKD-2 genes and that may be useful in the treatment of polycystic kidney disease. The assays of the invention comprise the contacting of genetically engineered cells expressing a mutant or truncated PKD gene product with a test agent and assaying for a decrease in the PKD mediated mutant phenotype. Characteristics associated with such a mutant phenotype include increased adherence to type I collagen coated surfaces; apical expression of NaK-ATPase on the cell membrane; increased expression of β -2-NaK-ATPase; and decreased focal adhesion kinase (FAK)

incorporation into focal adhesion complexes, and inability to form tubular structures in a gel matrix. To facilitate the screening methods of the invention, cells may be genetically engineered to express epitope tagged PKD gene products and/or epitope tagged PKD interacting proteins (PKD-IP). Such interacting proteins include, for example, focal

5 adhesion complex proteins such as FAK, paxillin, vinculin, talin and the like.

2. BACKGROUND OF THE INVENTION

Autosomal dominant polycystic kidney disease (ADPKD) is the most common lethal genetic disease inherited as a dominant trait in humans, with a prevalence of 1:1,000 live births. The disease afflicts approximately 6 million people world-wide,

10 and accounts for 5-7% of all patients on dialysis in the United States (Gabow, 1984, *Ann. Intern. Med.* **101**:238-247). Mutations in the *PKD-1* gene account for 85% of ADPKD, while mutations in the *PKD-2* gene account for 10% of the disease. In both cases, ADPKD is characterized by progressive, massive cystic enlargement of renal tubules resulting from increased proliferation, aberrant secretion, altered membrane protein

15 polarity, and extracellular matrix abnormalities correlated with a failure to down-regulate certain fetal genes after birth (Wilson, 1996, *In Polycystic Kidney Disease, Oxford Clinical Nephrology Series*, Watson M.L. and Torres V.E. eds. p. 124-163).

Approximately 50% of patients who inherit a mutant *PKD-1* gene will develop endstage renal failure, typically in the 5th decade of life, necessitating renal

20 replacement therapy by dialysis or transplantation. Since progression is usually slow and is a consequence of gradual loss of renal function as cysts continue to enlarge and destroy

intervening normal renal tubules, this presents a window of opportunity for potential drug therapies that would inhibit cyst expansion.

The *PKD-1* gene maps to human chromosome 16p13.3, has 46 exons, encodes a 14.5 kb transcript with a 12,912 basepair open reading frame and translates into a 4303 amino acid (≥ 462 kDa) protein, referred to as "polycystin-1" (European PKD Consortium, 1994, *Cell* **77**:882-894; International PKD Consortium, 1995, *Cell* **81**:289-298). The predicted amino acid sequence of the expressed protein suggests that the first 23 amino acids at the N-terminus act as a signal sequence, followed by two cysteine-flanked leucine rich repeats (LRR) which are strongly predictive of an extracellular location, protein-protein interactions and adhesion properties (Hughes et al., 1995, *Nat Genet* **10**:151-160). Other putative extracellular domains include a C-lectin-like motif and a region with high homology to the receptor for egg jelly of sea urchins (REJ), implying potential calcium influx regulation (Moy et al., 1996, *J. Cell. Biol.* **133**:809-817). The protein has several regions of high hydrophobicity and predicts 9-11 transmembrane domains and an intracellular carboxy terminal tail of 226 amino acids with putative binding sites for signal transduction molecules, including a SH2 site for tyrosine phosphorylation (YEMV) and two putative protein kinase C sites (RSSR) for serine phosphorylation.

The polycystin-1 protein is localized to areas of contact between the cell and matrix shortly after adhesion to type I collagen matrix. In addition, there is co-localization with defining focal adhesion proteins, namely $\alpha 2\beta 1$ -integrin, vinculin, α -actin, talin, paxillin, focal adhesion kinase (pp125^{FAK}) and pp60^{c-src} (Wilson et al., 1998, *J.*

Cell. Biol. Vol. **9**: 358A). Similar basally located polycystin-1-containing bodies have been demonstrated *in vivo* in human fetal ureteric bud epithelia in cell membrane regions in contact with type I collagen.

The overall predicted structure of the polycystin-1 protein and *in vitro* results with regard to matrix adhesion and phosphorylation assays, suggest that polycystin-1 functions as a matrix receptor mediating transfer of information from the extracellular matrix to the actin cytoskeleton, resulting in signal transduction that culminates in the nuclear regulation of gene transcription. This is suggested by the findings that all *PKD-1* mutations mapped to date would predict the translation of a truncated protein product, lacking varying amounts of the C-terminal domain (CTD), including a potential SH2 site, and mutations that result in the failure to down regulate fetal expression of the $\beta 2$ subunit of NaK-ATPase with consequent disruption of membrane polarization of NaK-ATPase (Peral et al., 1996, *Am. J. Hum. Gent.* **58**:86-96). Of additional interest, ADPKD epithelia have lower levels of PKD-1 tyrosine phosphorylation and fail to recruit FAK to the basally located multi-protein bodies.

In normal mature kidneys, NaK-ATPase is comprised of $\alpha 1\beta 1$ heterodimers located at the basolateral membranes of renal tubules and is associated with vectorial Na^+ export into the basal cell space (blood side) driving ion gradients for fluid reabsorption. In normal fetal kidneys and also in ADPKD kidneys, the $\beta 2$ subunit of NaK-ATPase heterodimerizes with $\alpha 1$ subunits which are “mis” targeted to apical plasma membranes, thus driving fluid secretion (Wilson et al., 1991, *Am. J. Physiol.* **260**:F420-

F430). The failure to repress $\beta 2$ transcription in adult kidneys results, therefore, in fluid secretion and expansion of renal tubule lumens into cysts.

Less is known about the *PKD-2* gene, which has been mapped to human chromosome 4q 21-23, encodes a 5.4kb transcript and translates into a predicted 110kDa protein, "polycystin-2" (Mochizuki et al., 1996, *Science* **272**:1339-1342). Unlike polycystin-1, the PKD-2 encoded protein has intracellular C- and NH- termini, 6 transmembrane domains with a putative EF hand and coiled-coil domain in the C-terminal region and putative SH3 sites in the N-terminal region. Yeast 2 hybrid studies have suggested potential interactions of *PKD-1* and *PKD-2* coiled-coil domains and some co-localizations have been suggested, however, in strict contrast to polycystin-1, polycystin-2 is not developmentally regulated (Tsoikas et al., 1997, *Proc. Natl. Acad. Sci.* **94**:6965-6970; Qian et al., 1997, *Nature Genet.* **16**:179-183).

3. SUMMARY OF THE INVENTION

The present invention relates to cell-based screening assays designed to identify agents that regulate the activity of polycystic kidney disease proteins (encoded by the PKD-1 and PKD-2 genes). The assay system of the invention is based on the use of eukaryotic cells containing naturally occurring mutations in the PKD genes and/or eukaryotic cells genetically engineered to express different forms of the polycystic kidney disease genes (PKD genes) including wild type, mutant, truncated, or epitope tagged PKD encoded proteins. In specific nonlimiting examples of the invention, human renal cells are engineered to express mutant or truncated forms of PKD protein and express a

mutant phenotype which includes one or more characteristics associated with renal epithelial cells from patients with polycystic kidney disease, for example, increased adherence to type I collagen coated surfaces; apical expression of NaK-ATPase on the cell membrane; increased expression of β -2-NaK-ATPase; and decreased FAK

5 incorporation into focal adhesion complexes, to name a few.

The assays provided by the present invention are designed to screen for compounds or compositions that modulate PKD activity, *i.e.*, compounds or compositions that act as agonists or antagonists of PKD, and thereby regulate PKD mediated signal transduction and extracellular matrix interactions with the plasma membrane and cell

10 cytoskeleton. The invention also relates to assays designed to screen for compounds or compositions that modulate *PKD* gene expression, *i.e.*, compounds that modulate expression of *PKD*, or production or activity of transcription factors involved in *PKD* gene expression. Agents identified using the cell-based assays of the invention may be useful in the treatment of renal cystic disease.

15 In addition, the assays of the invention may be used to identify and/or validate protein interactions between PKD and binding partners within the cell, identified using different model host systems, *i.e.*, *C.elegans*. To this end, cells endogenously or genetically engineered to express PKD may be further engineered to express PKD interacting proteins. Various standard assays may be used to validate such protein

20 interactions and/or to serve as a basis for screening assays designed to identify agents that regulate the formation of such protein complexes.

3.1 DEFINITIONS

As used herein, italicizing the name of a gene shall indicate the gene, in contrast to its encoded protein product which is indicated by the absence of italicizing.

For example, *PKD* shall mean the *PKD* gene, whereas PKD shall indicate the PKD

5 encoded protein product. Unless otherwise specified, *PKD* shall refer to a *PKD-1* or *PKD-2* gene of any species, whereas PKD shall indicate an PKD-1 or PKD-2 encoded protein products (also referred to as polycystin-1 and polycystin-2, respectively) of any species. Such *PKD* genes and encoded proteins include, but are not limited to, those described by Mochizuki T et al. (1996, *Science* **272**:1339-1342); The International
10 Polycystic Kidney Disease Consortium (1995, *Cell* **81**:289-298); and Barr and Sternberg (1999, *Nature* **401**:386-389).

4. BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1A-B. Characterization of anti-polycystin-1 antibody. Western immunoblot analysis using 1:12,500 dilution of anti-peptide (aa 4161-4191) polyclonal
15 antiserum raised in rabbits. Figure 1A. PKD1-C-terminal domain (aa 4105-4303) fusion protein before (lane 1) and after (lane 2) IPTG induction. Human fusion protein is denoted by * the predicted molecular weight. Incubation with pre-immune serum did not detect bands. Figure 1B. Human fetal collecting tubule cell lysates were incubated in immune serum (lanes 1 and 2) or pre-immune serum (lanes 3 and 4). A >440kDa band
20 was detected in lanes 1 and 2.

Figure 2A-H. Peptide competition of immunocytochemical staining with anti-polycystin-1 antibody. Panels A-D, immunohistochemistry without pre-adsorption; Panels E-H, immunohistochemistry after preadsorption with 31 amino acid polycystin-1 peptide (100 μ g/ml). A and E, 16-week human fetal kidney (anti-polycystin-1 1:500); B and F, 39 year normal adult kidney medulla (anti-polycystin-1 1:500); C and G, endstage autosomal dominant polycystic kidney disease (anti-polycystin-1 1:500); D and H, cultured autosomal dominant polycystic disease epithelia (anti-polycystin-1 1:1,000).

Figure 3A-H. Localization of polycystin-1 in human renal tissues and epithelial cell lines. Immunocytochemical localization of polycystin-1 in human renal tissues (Panels A-D) and nephron segment-specific epithelial cell cultures (Panels E-H). A. Human fetal kidney (12 weeks gestation). Note staining in ureteric bud-derived collecting duct epithelia (arrows). X 125. B. Normal, perfused, adult human kidney: inner medullary collecting tubules show light staining. X 60. C. Cystic ADPKD kidney: intense staining in cyst lining epithelia. X 60. D. Absence of staining in ADPKD kidney incubated with pre-immune serum. X 100. E. Normal human fetal collecting tubule epithelial cell line: intense staining. X 125. F. Normal adult collecting tubule cell line: weak staining. X 60. G. ADPKD epithelial cell line: Intense staining. X 125. H. Absence of staining in ADPKD epithelial cell line incubated in preimmune serum. X 100.

Figure 4A-C. Co-immunoprecipitation of polycystin-1 with focal adhesion proteins. Figure 4A. Specificity of polycystin-1 immunoprecipitation in human fetal

collecting tubule cells. Lane 1, immunoprecipitation with anti-polycystin-1 followed by Western analysis with anti-polycystin 1. Note major band at >440kDa and minor band at 210kda. Lane 2, immunoprecipitation control using protein A/G agarose beads omitting anti-polycystin-1 antiserum. Lane 3, control omitting anti-polycystin-1 antiserum in immunoblot. Lane 4, immunoprecipitation control using preimmune serum instead of anti-polycystin-1 antiserum. Figure 4B. Co-immunoprecipitation with polycystin-1 in normal human fetal collecting tubule epithelia. Immunoprecipitation with anti-polycystin-1 followed by Western immunoblot with anti-polycystin-1 (Lane 1); anti-vinculin (lane 2); anti-paxillin (lane 3); anti- pp125^{FAK} (lane 4). Note the presence of 130kDa vinculin band, 68kDa paxillin band and 125kDa FAK band in lanes 2, 3 and 4, respectively. Equal protein loading of 20 mg per lane. C. Co-immunoprecipitation of proteins with polycystin-1 in ADPKD epithelia. Conditions as in B. Note presence of vinculin (lane 2) and paxillin (lane 3) in the co immunoprecipitates but the absence of pp125^{FAK} (lane 4).

Figure 5A-B. Western Immunoblot analysis of polycystin-1 in human renal epithelia anti-C-terminal peptide antibody 1:12,500. Figure 5A. Tissue extracts SDS-PAGE: Lanes 1 and 2 fetal kidneys; lanes 3 and 4 normal adult kidneys; lanes 5 and 6 ADPKD kidneys. Equal protein loading 20 mg per lane. Figure 5B. Cultured epithelial cell extracts, SDS-PAGE: Lanes 1 primary culture of human fetal collecting tubule epithelia (HFCT); lane 2 immortalized HFCT clonal cell line; lane 3, primary culture of ADPKD epithelia; lane 4, immortalized ADPKD clonal cell line. Equal protein loading 20 mg per lane. Note band of >440 kDa ferritin marker position, consistent with the

predicted molecular weight of the full length polycystin-1 protein. Additional band at 210kDa was sometimes also detected.

Figure 6A-C. Adhesion of normal and ADPKD epithelia to extracellular matrix proteins. Figure 6A. Differential adhesion of human renal primary epithelial cell cultures derived from different nephron segments on type I collagen, (1000 cells, 48 hr.) in serum-free media. Lane 1, human fetal proximal tubule; lane 2, human fetal collecting tubule; lane 3, normal human adult proximal tubule; lane 4, normal adult thick ascending limb of Henle's loop; lane 5, normal adult collecting tubule; lane 6, autosomal dominant polycystic kidney disease cyst epithelia (ADPKD). * $p < 0.01$. Figure 6B. Differential adhesion (2,000 cells 12 hr) of primary cultures of ADPKD and normal adult thick ascending limb (NHTAL) epithelia on different matrix proteins. * $p < 0.01$ ADPKD versus NHTAL on corresponding matrix. Figure 6C. Comparative cell-type-dependent adhesion to type I collagen of human immortalized cell lines after differentiation at non-permissive temperature 37°C for 9 days, versus primary cultures. Lane 1, human fetal proximal tubule (HF PT) primary culture, passage 1; lane 2, HF PT immortalized, differentiated clonal line; lane 3, normal human fetal collecting tubule (HFCT) immortalized, differentiated clone; lane 4, immortalized differentiated NH PT clone; lane 5, ADPKD primary epithelial culture, P1; lanes 6, 7, and 8, immortalized differentiated ADPKD independent clones: 4,000 cells, 4 hr. Data are expressed as means \pm SEM. * $p < 0.01$.

Figure 7A-C. $\alpha 2\beta 1$ integrin and adhesion of ADPKD epithelia to type I collagen. Figure 7A. Immunocytochemistry of $\alpha 2\beta 1$ -integrin (1: 250, Chemicon) in

ADPKD epithelia, 48 hr. adhesion to type I collagen. Note extensive fibrillar staining at apparent points of focal contact with the substratum. Figure 7B. Western immunoblot analysis of $\alpha 2$ -integrin content of different cell types, 48 hr. adhesion to type I collagen. Lanes 1 and 2 HF PST; lanes 3 and 4 HF CT; lanes 5 and 6, NH PST; lane 7, NH TAL; lanes 8 and 9 ADPKD. * anti-actin to demonstrate equal protein loading, 20 mg per lane, equal exposure of blot. Figure 7C. Differential effects of the $\alpha 2\beta 1$ -integrin blocking antibody, 6F1, on cell-type-dependent adhesion (4 hr.): incubation in the presence of 10 mg/ml monoclonal anti- $\alpha 2\beta 1$ -integrin blocking antibody, 6F1 (filled bars) or irrelevant monoclonal antibody, PCNA (Signet, Deedham, MA) (open bars). Values expressed as mean \pm SEM. * $p < 0.05$ by comparison to its pair. Adherence in the presence of 1% fetal bovine serum was similar to that in the presence of anti-PCNA. Lanes 1 and 2 HF PT; lanes 3 and 4 NH PST; lanes 5 and 6 NH CT; lanes 7-12 ADPKD, 3 independent kindreds.

Figure 8A-C. Immunoblot analysis of focal adhesion proteins in cultured human renal tubule epithelia. Figure 8A. Anti-vinculin monoclonal 1:7,500 (Sigma): 20 mg total protein per lane. Lane 1, normal adult proximal tubule epithelia; lane 2, normal adult thick ascending limb epithelia; lane 3, normal adult collecting tubule epithelia; lane 4, early stage ADPKD; lane 5, endstage ADPKD. Equal protein loading 20 mg per lane. Note increased content of vinculin in ADPKD epithelia. Figure 8B. Anti-paxillin monoclonal, 1:10,000 (Transduction Laboratories). 20 mg total protein per lane. Lanes as in A. Note increased content of paxillin in ADPKD epithelia. Figure *C.

Anti-pp125^{FAK} monoclonal, 1:10,000 (Transduction Laboratories). 20 mg total protein per lane. Lanes as in A. Note decreased content of pp125^{FAK} in ADPKD epithelia.

Figure 9A-I. Co-localization of polycystin-1 with integrins and associated proteins in ADPKD epithelia after short-term adherence to type I collagen. Confocal microscopy of double immunofluorescent labeling of ADPKD epithelial cells 4 hr. after adhesion to type I collagen. All cells were stained after incubation with 1:500 affinity purified, polyclonal anti-peptide C terminal polycystin-1 antibody (Panels B, E and H) followed by incubation with one of the following monoclonal antibodies: anti α 2 β 1-integrin antibody, 1:100, (Chemicon), (panel A); anti-vinculin, 1:500, Sigma (panel D) or anti-paxillin, 1:250, (Sigma) (panel G). Monoclonal antibodies were visualized by coupling to FITC-labeled anti-mouse IgG and polyclonal antibody localization was visualized by rhodamine-coupled anti-rabbit IgG. Merging of the images (panels C, F and I) and resultant yellow/orange demonstrate co-localization of polycystin-1 with α 2 β 1-integrin (C); vinculin (F) and paxillin (I). Plane of focus was at the level of contact of the cell with the substratum. X 250.

Figure 10A-F. Co-localization of polycystin-1 with integrin and associated proteins in normal human fetal collecting tubule epithelia after short-term adherence to type I collagen. Confocal microscopy of double immunofluorescent labeling of fetal collecting tubule epithelial cells 4 hr. after adhesion to type I collagen. Panel A, anti- α 2 β 1-integrin; Panels B and E, anti-polycystin-1; Panel D, anti-vinculin; Panels C and F, merged images.

Figure 11A-C. Tyrosine phosphorylation of polycystin-1. Figure 11A.

Immunoblot analysis of polycystin-1 (1:12,500) in proliferating, adherent cultured ADPKD and normal epithelial cells. Arrow denotes position of 209kDa marker. Note major band at >400kDa and a single additional band of varying intensity at 225kDa.

- 5 Lane 1, ADPKD primary; lane 2, ADPKD immortalized clone; lane 3, NHCT primary; lane 4, NH TAL primary; lane 5, HF CT conditionally immortalized clone; lane 6, HFCT primary; lane 7, HFPT conditionally immortalized clone. Equal protein loading 20 mg per lane. Figure 11B. Identical immunoblot as in A but stripped and reprobed with anti-phosphotyrosine antibody (RC-20, 1:2,500, Transduction Labs). Note
- 10 immunoreactivity in the ≥ 440 kDa and 225 kDa bands, together with additional, lower molecular weight proteins. (Arrow shows position of 209 kDa marker). Figure 11C. Immunoprecipitation of confluent ADPKD and normal epithelial cells with anti-phosphotyrosine antibody (RC-20) followed by Western immunoblot analysis with anti-polycystin-1 antibody shows a major ≥ 440 kDa band, attesting to the
- 15 tyrosine-phosphorylation of polycystin-1. Lane 1, ADPKD primary epithelia; lane 2, NHCT primary epithelia; lane 3, HFCT primary epithelia.

Figure 12A-C. Depictions of proposed structures of PKD-1 and PKD-2.

Figure 12A. Schematic representation of PKD-1 encoded gene product (Wilson et al., 1999, Exp Nephrol 7:114-124). Figure 12B. Schematic representation of PKD-2 encoded

20 gene product (Wilson and Guay-Woodford 1999, Seminars in Nephrology 19: 123-132)

Figure 12C. Model of polycystin-1 in a multi-protein complex at the cell membrane (Wilson et al., 1999, *Laboratory Investigation* 79:1-13).

Figure 13. PKD-1-epitope tagged eukaryotic expression vectors.

5. DETAILED DESCRIPTION OF THE INVENTION

The present invention relates to cell-based screening assays designed to identify agents that regulate the activity of polycystic kidney disease proteins (encoded by the PKD-1 and PKD-2 genes). The assay system of the invention is based on the use host cells that naturally express different forms of the polycystic kidney disease genes (PKD1 and PKD-2 and/or host cells genetically engineered to express different forms of the polycystic kidney disease genes (PKD1 and PKD-2) including wild type, mutant, truncated, or epitope tagged PKD proteins. In specific nonlimiting examples of the invention, human renal cells are engineered to express mutant or truncated forms of PKD protein and express a mutant phenotype which includes one or more characteristics associated with renal epithelial cells from patients with polycystic kidney disease, for example, increased adherence to type I collagen coated surfaces; apical expression of NaK-ATPase on the cell membrane; increased expression of β -2-NaK-ATPase; and decreased FAK incorporation into focal adhesion complexes, to name a few.

5.1 GENERATION OF RECOMBINANTLY ENGINEERED PKD EXPRESSING CELLS

In accordance with the invention, a cell-based assay system can be used to screen for agents that modulate the activity of PKD, and thereby modulate the mutant phenotype associated with polycystic kidney disease. To this end, host cells, such as

human renal epithelial cells, are genetically engineered to express wild type, mutant, truncated or epitope tagged *PKD* proteins for use in the screening assays of the invention.

The cloning and characterization of the *PKD-1* and *PKD-2* genes from various species has been described (Mochizuki T et al., 1996, *Science* **272**:1339-1342; 5 The International Polycystic Kidney Disease Consortium, 1995, *Cell* **81**:289-298; and Barr and Sternberg, 1999, *Nature* **401**:386-389). In addition, mutant forms of the *PKD* receptor have been identified and described (European Polycystic Kidney Disease Consortium, 1993, *Cell* **75**:1305-1315; Peral et al., 1996, *Am J. Hum Genet* **58**:86-96). For purposes of the present invention, host cells are genetically engineered to express any 10 of the different forms of the *PKD* genes so long as the result is the expression of a mutant phenotype. Indeed, in some instances, over expression of wild -type *PKD* protein may result in expression of a mutant phenotype.

The nucleotide sequence coding for a *PKD* protein (*PKD-1* or *PKD-2*), or fragment or other derivative thereof, can be inserted into an appropriate expression 15 vector, *i.e.*, a vector which contains the necessary elements for the transcription and translation of the inserted protein-coding sequence. In specific embodiments, the nucleotide sequence encoding a wild type or mutant *PKD* gene is cloned into an appropriate expression vector. Alternatively, nucleotide sequences encoding a functionally active domain of *PKD* can be cloned into an expression vector. For 20 example, a fragment of *PKD* comprising functional domains such as an extracellular, transmembrane or intracellular C-terminal domain of the *PKD* protein can be expressed within a cell. In a specific embodiment, nucleotide sequences encoding the C-terminal

domain of PKD, *i.e.*, the C-terminal 226 and 200 amino acids of PKD-1 and PKD-2, respectively, are cloned into an appropriate expression vector. Such vectors may further comprise sequences that result in a physical association between the C-terminus of PKD and the plasma membrane. In a first nonlimiting embodiment, a myristylation sequence
 5 may be linked to the PKD encoding sequences to ensure targeting to the plasma membrane. Such myristylation sequences include the N-terminal myristylation sequence from pp60 src (Kaplan et al., 1988, *Mol Cell Biol* **8**:2435-41). Further, a leader sequence, such as an N-terminal pre-protrypsin leader sequence, may be included to ensure membrane insertion of a truncated PKD protein.

10 In addition, PKD fusion proteins may be engineered to include a short stretch of residues corresponding to an epitope or other detectable label to facilitate subsequent biochemical and/or immunological analysis (*e.g.*, “epitope tagging”). This is achieved by linking the sequence of the detectable tag to the PKD coding sequence. Suitable epitope tags include but are not limited to EGFP, FLAG, HA, 6-HIS, AU1, and
 15 green fluorescent protein (GFP) tags (see, Ausubel et al. (eds.), 1993, *Current Protocols in Molecular Biology*, John Wiley & Sons, N.Y.). An example of another detectable tag is green fluorescent protein.

Vectors encoding PKD proteins can be plasmid, viral or other vectors known in the art, for replication and expression in cells, including but are not limited to
 20 eukaryotic cells. Expression of a nucleic acid sequence encoding a PKD protein or peptide fragment may be regulated by any promoter/enhancer element known in the art to act in eukaryotic, preferably mammalian cells. Such promoters which may be used to

control PKD expression can be inducible or constitutive. Such promoters include, but not limited to, the SV40 early promoter region (Bernoist and Chambon, 1981, *Nature* **290**:304-310), the promoter contained in the 3' long terminal repeat of Rous sarcoma virus (Yamamoto, et al., 1980, *Cell* **22**:787-797), the herpes thymidine kinase promoter (Wagner et al., 1981, *Proc. Natl. Acad. Sci. U.S.A.* **78**:1441-1445), the regulatory sequences of the metallothionein gene (Brinster et al., 1982, *Nature* **296**:39-42); prokaryotic expression vectors such as the beta-lactamase promoter (Villa-Kamaroff, et al., 1978, *Proc. Natl. Acad. Sci. U.S.A.* **75**:3727-3731), or the *tac* promoter (DeBoer, et al., 1983, *Proc. Natl. Acad. Sci. U.S.A.* **80**:21-25); promoter elements from yeast or other fungi such as the Gal 4 promoter, the ADC (alcohol dehydrogenase) promoter, PGK (phosphoglycerol kinase) promoter, alkaline phosphatase promoter. In addition, transcriptional control regions, which exhibit tissue specificity such as, for example, control regions which are active in renal epithelial cells may be used. The necessary transcriptional and translational signals can also be supplied by the native PKD gene and/or its flanking region.

The expression of the PKD protein may be regulated at the transcriptional level through the use of tightly control promoter systems that allow for inducible expression of the PKD proteins. Such promoters include, for example, those utilized in a tetracycline-controlled system (Gossen et al., 1993, *Trends Biochem Sci* **18**:471-475; Barinaga, 1994, *Science* **265**:26-28; Damke et al., *Methods in Enzymol.* **257**:209-220).

Vectors to be used for expression of PKD may optionally comprise one or more selectable markers (e.g., an antibiotic resistance gene). Such vectors can remain

episomal or become chromosomally integrated, as long as they can be transcribed to produce the desired cell lines expressing PKD.

Any of the methods well known to those skilled in the art can be used for the insertion of DNA fragments to construct expression vectors containing appropriate transcriptional/translational control signals and the PKD protein coding sequences. The methods to be used for construction of such vectors may include *in vitro* recombinant DNA and synthetic techniques and *in vivo* recombinants. Methods commonly known in the art of recombinant DNA technology which can be used are described in Ausubel et al. (eds.), 1993, *Current Protocols in Molecular Biology*, John Wiley & Sons, N.Y.

The present invention encompasses cell-based screening assays that utilize cell lines recombinantly engineered to express PKD proteins. In a preferred embodiment of the invention mammalian cells, such as primary or immortalized cultures of renal epithelial cells, are transfected with the recombinant vectors engineered to express PKD protein. Human renal cell lines may be derived from normal human adult kidney proximal tubules, normal human adult kidney collecting tubules, normal human adult kidney thick ascending limb tubules, normal human fetal kidney proximal tubules, normal fetal adult kidney collecting tubules, human adult autosomal dominant polycystic kidney disease epithelia and human adult autosomal recessive polycystic kidney disease epithelia. Such renal epithelial cells may be cultured in serum-free, cell-type specific media as described in Wilson et al. (1985, *Am. J. Physiol.* **248**:F436-F443; and 1986, *Kidney Int.* **30**:371-378) and immortalized as described in Racusen et al., (1995, *Kidney Int.* **48**:536-543).

In another embodiments of the invention, the screening assays may be utilize non-human cells that contain a homolog of human *PKD-1* or *PKD-2* such as cells of *Caenorhabditis elegans*.

Various methods commonly known in the art for delivery of recombinant DNA to cells in tissue culture may be used. Such methods include electroporation, lipofection, calcium phosphate mediated transfection, or viral infection. Usually, the method of transfer includes the transfer of a selectable marker to the cells. The cells are then placed under selection to isolate those cells that have taken up and are expressing the transferred gene.

In addition to expression of introduced PKD proteins, host cells may be genetically engineered to express proteins known to, or suspected of, interacting with PKD proteins expressed within the cell. Such proteins include focal adhesion complex proteins such as FAK, paxillin, vinculin, talin, PKD-2 and *C. elegans* proteins such as unc-54 and act-5 and their human homologs, and the like. The tagging of such proteins will facilitate biochemical and immunological analysis of protein interactions for use in the screening assays of the invention. To this end, cells that endogenously express wild type or mutant PKD genes may be engineered to express epitope tagged PKD interacting proteins. Alternatively, cells may be co-engineered to express the various different forms of the PKD protein, as described *supra*, and epitope tagged PKD interacting proteins. Cells expressing such tagged proteins provide a tool for cell-based screening assays designed to identify agents capable of regulating, *i.e.*, inhibiting or altering the affinity of, PKD protein interactions, and modifying the cellular phenotype. As a specific example, a

human renal epithelial cell may be cotransfected with (i) a first gene encoding an epitope-tagged PKD-1 comprising the C-terminus and a membrane localization sequence and (ii) a second gene encoding an epitope tagged protein that interacts with PKD-1 at the cell membrane. The interaction of the proteins at the membrane may be visualized by immunological staining techniques. Such cells may be used to screen for agents that block the interaction of the proteins.

In a specific embodiment of the invention, several PKD-1-epitope tagged eukaryotic expression vectors were constructed and shown not only to transfect with high efficiency into normal renal epithelia and 293 cells, but also to be localized into basally located bodies at the cell membrane. The constructs include EGFP, FLAG and/or 6-His tags of the cytoplasmic portion of the C-terminal domain of PKD, with or without a myristylation sequence to ensure targeting to the plasma membrane. Also, similarly tagged constructs containing up to 5 transmembrane domains as well as the C-terminal domain (Figure 13) were constructed and expressed in mammalian renal epithelia and 293 cells.

In addition to myristylation signals, membrane insertion of the FLAG tagged PKD-1 constructs was achieved via use of an N-terminal preprotrypsin leader sequence. Membrane insertion of the epitope tagged PKD-1 proteins, predicted to have 5 transmembrane (TM) domains and the CTD, was confirmed by immunohistochemistry. In contrast, the pEGFP/PKD-1 constructs which had no leader sequence were expressed as cytoplasmic proteins. The myristylated constructs appear to have the most efficient membrane localization to the basally located bodies of all of the constructs so far tested.

5.2. ASSAY SYSTEMS

In utilizing the cell-based assay systems of the invention, cells expressing PKD proteins are exposed to a test compound or to vehicle controls (*e.g.*, placebos).

After exposure, the cells can be assayed to measure the expression and/or activity of the

5 components of the signal transduction pathway of the PKD receptor. For example, cells can be assayed for alterations in characteristics associated with renal cells from patients suffering from polycystic kidney disease such as adherence to type I collagen coated surfaces; expression of NaK-ATPase on the cell membrane; expression of β -2-NaK-ATPase; and FAK incorporation into focal adhesion complexes, proliferation, the

10 inability to form tubular structures in a gel matrix and the expression of various fetal genes. In many instances, standard immunoassays which include Western blots, radioimmunoassays, ELISAs, immunoprecipitation assays and fluorescent immunoassays, to name but a few, may be utilized to detect changes in localization of PKD proteins and/or changes in interactions with other cellular proteins. In preferred

15 embodiments, the assays are performed in 96-well plates to enable high-throughput screening and 96-well based scintillation counting instruments.

To confirm that any observed alterations in phenotype are due to a specific effect of the test compound on mutant PKD signal transduction, the test compound is also tested for its ability to alter the phenotype of cells expressing wild type

20 PKD.

The cell based assays are designed to identify compounds which affect *PKD-1* activity. Compounds that may affect *PKD-1* activity include but are not limited to compounds that bind to the *PKD-1* functional domains and either activate signal transduction (agonists) or block activation (antagonists). Compounds that affect *PKD-1* activity by affecting *PKD-1* gene expression, including molecules that affect transcription can also be identified using the screens of the invention. However, it should be noted that the assays described can also identify compounds that modulate *PKD-1* signal transduction (e.g., compounds which affect downstream signaling events and participate in transducing the PKD-mediated signal). The identification and use of such compounds which affect signaling events downstream of *PKD-1* and thus modulate effects of *PKD-1* on the development of polycystic kidney disease are within the scope of the invention.

The compounds which may be screened in accordance with the invention include, but are not limited to proteins peptides, antibodies and fragments thereof, and other organic compounds (including, but not limited to, peptidomimetics) that bind to *PKD-1* and either activate the activity of *PKD-1* (i.e., agonists) or inhibit the activity of *PKD-1* (i.e., antagonists). Compounds may include, but are not limited to, peptides such as, for example, soluble peptides, including but not limited to members of random peptide libraries; (see, e.g., Lam, K.S. et al., 1991, *Nature* **354**:82-84; Houghten, R. et al., 1991, *Nature* **354**:84-86), and combinatorial chemistry-derived molecular library made of D- and/or L- configuration amino acids, phosphopeptides (including, but not limited to, members of random or partially degenerate directed phosphopeptide libraries; see, e.g.,

Songyang, Z. et. al., 1993, *Cell* **72**:767-778). Other agents, not related to peptides, may also be screened, according to the invention.

To assay for a test compound's ability to regulate PKD activity, adherence to type I collagen coated surfaces may be assayed using tissue culture surfaces coated with type I collagen. Cells expressing mutant PKD have a 2- to 3-fold increase in adhesion to type I collagen coated surfaces as compared to wild type cells. Thus, the ability of a test compound to decrease levels of mutant cell adherence to type I collagen coated surfaces, below those levels seen with cells treated with a vehicle control, indicates that the test agent inhibits the signal transduction mediated by PKD.

To assay for cell adherence, cells may be plated onto 96-well cluster plates pre-coated with type I collagen, for a time sufficient to allow attachment in the presence of either a test agent or a vehicle control. Following attachment, the non-attached cells are removed by aspiration. Measurement of cell number attached to each plate is then carried out. For example, a colorimetric assay can be utilized to measure cell number attached to each well in the plate, read and quantitated using a microplate reader (Mossmann, 1983, *Imm Meth* **65**:55-63).

Increased adhesion may also be determined using assays designed to detect the level of binding of cells to one another to form cluster of cells. The ability of a test agent to decrease the level of cell adherence, below those levels seen with cells treated with a vehicle control, indicates that the test agent inhibits the signal transduction mediated by PKD.

The ability of cells to form tubular structures may be evaluated by seeding the cells plus test agent in Matrigel or a comparable gel culture system.

Assays for expression of NaK-ATPase on the cell membrane may also be used in conjunction with genetically engineered cells expressing PKD for identification of agents capable of regulating PKD activity. Since PKD mutant cells express NaK-ATPase on their apical cell membranes, whereas normal renal epithelia have exclusively basal membrane located NaK-ATPase, this phenotype can be used to assess the ability of a test agent to reverse NaK-ATPase apical localization. Expression of NaK-ATPase in the cell membrane can be measured in the presence of a test compound or vehicle control, using for example, a fluorescent antibody against the NaK-ATPase sodium pump. Polarity of fluorescence may be easily assessed on a fluorescent reader. The ability of a test agent to decrease the level of NaK-ATPase expression on the apical cell membrane, below those levels seen with cells treated with a vehicle control, indicates that the test agent inhibits the signal transduction mediated by PKD.

In yet another embodiment of the invention, the ability of a test agent to decrease the expression of the $\beta 2$ subunit of NaK-ATPase may be assayed to identify regulators of PKD activity. This assay is based on the observation that cells expressing mutant PKD express β -2 NaK-ATPase, while normal adult renal epithelial cells do not. Recombinantly engineered cells may be grown to confluence in 96-well cluster plates and the expression of the $\beta 2$ subunit examined using a fluorescent detection system in the presence of a test agent or vehicle control. The ability of a test agent to decrease the level of $\beta 2$ subunit of NaK-ATPase expression, below those levels seen with cells treated with

a vehicle control, indicates that the test agent inhibits the signal transduction mediated by mutant PKD.

The incorporation of PKD proteins into focal adhesion clusters can also be used to assess the ability of a test compound to regulate the activity of mutant PKD.

5 Other proteins known to localize to focal adhesion clusters include $\alpha 2 \beta 1$ integrin, vinculin, talin and paxillin. To this end, focal adhesion complex proteins, such as paxillin, vinculin and talin, may be genetically engineered so as to be labeled with epitope tags. Antibodies immunoreactive against the epitope tags may be used in the assays of the invention to determine whether a test compound is capable of altering the
10 protein composition of focal adhesion complexes. For example, double labeling immunofluorescent studies may be conducted to determine whether PKD co-localizes in the cell with $\alpha 2 \beta 1$ -integrin, vinculin and paxillin using fluorescent labeled antibodies immunoreactive against each of the proteins. Numbers of focal adhesion clusters can be assessed by reading on a fluorescent plate reader and the location of focal clusters can be
15 assessed microscopically.

In normal cells, focal adhesion kinase (FAK) is incorporated into focal adhesion complexes, while in mutant cells FAK fails to be incorporated into such complexes. Thus, agents may be screened for their ability to increase the level of FAK incorporated into focal adhesion complexes. To this end, FAK may be genetically
20 engineered so as to be labeled with epitope tags such as FLAG, HA, green fluorescent protein or 6-His. Antibodies immunoreactive against the epitope tags may be used in the assays of the invention to determine whether a compound is capable of altering the

localization of FAK in focal adhesion complexes. For example, labeling immunofluorescent studies may be conducted to determine whether FAK localizes to focal adhesion complexes.

5.3. USE OF *C.ELEGANS* MODEL SYSTEM

Compounds identified via cell based assays such as those described herein may be useful, for example, in elaborating the function of PKD gene products and for ameliorating the symptoms associated with polycystic kidney disease. Assays for testing the efficacy of compounds identified in the cell-based assays can also be tested in model systems for polycystic kidney disease based on other types of organisms. Such model systems may be used as test substrates for the identification of compounds that may be effective in treating polycystic kidney disease.

To this end, *C. elegans* can be used as a model system. The *C. elegans* genome has homologs of both the human *PKD-1* and *PKD-2* genes. Thus, the *C. elegans* ~~excretory cell~~ provides a new model system for the study of essential genes (including the polycystic kidney disease genes *PKD-1* and *PKD-2*) in the development and maintenance of the renal epithelia of the mammalian nephron. An understanding of the effects of *PKD-1* and *PKD-2* mutations in *C. elegans* will provide additional insights into the function of these genes and form the basis for a screening approach for the validation of compounds identified in the cell based assays with potential therapeutic activity in the treatment of human polycystic kidney disease. For example, a mutated form of *PKD-1* or *PKD-2* may be introduced into *C. elegans* cells or organisms and its effect on

excretory cell function and cell attachment bodies evaluated. Such genetically modified *C. elegans* cells or organisms may be used to screen for agents that reverse the effects of the mutation.

In addition, use of a yeast two-hybrid screen may be used for the

5 identification of *C.elegans* PKD-1 and PKD-2 interacting proteins. To confirm that the interacting proteins from *C.elegans* reflect a molecular interaction that parallels that of PKD proteins in human cells, full length cDNA clones representing the human homologs of the identified PKD interacting proteins may be obtained, sequenced and subcloned into a suitable expression vector. Antisera may be prepared against selected peptides derived

10 from the open reading frames of the obtained cDNAs. To validate PKD-1 and PKD-2 protein interactions identified in the *C. elegans* yeast two-hybrid screen, renal cells engineered to express tagged human homologs of the PKD-1 and PKD-2 interacting proteins may be used to determine whether there is co-localization with PKD proteins.

To this end, renal epithelial cells engineered to express tagged PKD

15 polypeptides may be used to determine whether the putative interacting proteins colocalize with the PKD protein products. Such studies may be conducted using standard immunoassays which include western blots, radioimmunoassays, ELISAs, immunoprecipitation assays and fluorescent immunoassay, to name but a few. For example, co-immunoprecipitation of PKD interacting proteins and PKD proteins can be

20 demonstrated using PKD-1 and PKD-2 antisera in selected human renal epithelial cell lines engineered to express PKD-1 and PKD-2 proteins. The selected cells may also be transfected with epitope-tagged putative PKD interacting proteins. Additionally, PKD

immunoprecipitations followed by Western immunoblotting using antisera raised against PKD interacting proteins in non-transfected renal cells may be performed to determine in protein interactions occur at physiological levels of protein expression. Identification of such specific protein interactions provides a target for drug screening.

5 *C. elegans* proteins that interact with PKD proteins in a manner that parallels PKD interactions in human renal cells may be used as a basis for a primary screen, in *C. elegans*, to identify agents that modulate the protein interaction. The ability of modulating agents identified in the *C. elegans* screen to modulate the interacting human homologs may then be confirmed using human renal cell lines transfected
10 according to the invention.

6. EXAMPLE: EXPRESSION OF *PKD-1* IN HUMAN TISSUES AND RENAL EPITHELIAL CELLS

 The expression of the *PKD-1* encoded protein “polycystin-1” in human
15 tissues and renal epithelial cell lines and the mechanisms of adhesion of polycystin-1-expressing renal epithelial cells to matrix proteins was examined. *PKD-1* mRNA is highly expressed in normal human fetal kidneys in the ureteric bud epithelium, as well as in the cystic epithelial cells of ADPKD kidneys (Wilson, 1997, *Am J Physiol* **272**:F434-F442; Ward et al. 1996, *Proc Natl Acad Sci USA* **93**:1524-1528; Geng et al.
20 1996, *J Clin Invest* **98**:2674-2682). Previously devised techniques for generating pure cell lines of different human renal tubule epithelial cell types from normal adult and fetal kidneys, as well as from ADPKD cysts have been described (Wilson et al., 1985, *Am J*

Physiol **248**:F436-F443; Racusen et al., 1995, *Kid Int* **48**:536-543; Cuppage et al., 1980, *Kid Int* **17**:372-381). For the present studies, polycystin-1-expressing normal and ADPKD epithelial cell lines were used to study the biological function of polycystin-1 in response to matrix adhesion.

5

6.1. MATERIALS AND METHODS

6.1.1 TISSUE CULTURE

Primary and immortalized cells of normal fetal and adult proximal tubules, thick ascending limbs of Henle and collecting tubules and of ADPKD cyst epithelia were cultured in serum-free, cell-type specific media as described previously (Wilson et al., 1985, *Am J Physiol* **248**:F436-F443, Wilson et al., 1986, *Kidney Int* **30**:371-378 ; Racusen et al., 1995, *Kidney Int* **48**:536-543). Cells from each cell type were immortalized by retroviral transduction of the SV40 tsA58 U19 vector and selection in G418 at 33 °C for 6 weeks. Clonal, immortalized ADPKD cell lines have a truncating mutation in exon 26 (Harris, P.C., personal communication). Prior to experimentation all immortalized cell lines were cultured for 7-14 days at 37 °C, to terminate proliferation and maximize differentiation.

6.1.2 RECOMBINANT FUSION PROTEIN PRODUCTION

A fusion protein containing the C-terminal of human polycystin-1 (amino acids 4105-4303) was amplified by PCR from a 2kb PKD1 cDNA clone obtained by

RT-PCR and subcloned into PET-32 LIC (Novagen). The fusion protein was expressed in BL-21 (D3) plys S competent cells and purified using S-tag affinity chromatography.

6.1.3. IMMUNOCYTOCHEMISTRY

The anti-polycystin-1 antibody was raised in rabbits against a purified
 5 synthetic 31 amino acid peptide corresponding to amino acids 4161-4191 in the predicted
 intracellular portion of polycystin-1, proximal to the C-terminal: sequence
 LPSRSSRGSKVSPDVPPPSAGSDASHPSTSS. Antiserum specificity was confirmed
 by Elisa, immunoblot and immunocytochemical analyses, before and after affinity
 purification; by lack of staining with pre-immune sera and competition of
 10 immunoreaction by preadsorption with the appropriate peptide. Immunocytochemistry
 was carried out using an avidin-biotin-peroxidase system (Vectastain, Vector
 Laboratories) and aminoethylcarbazole as chromogen 9 (red color). Staining patterns
 were identical when carried out on frozen and paraformaldehyde (4%)-fixed material.
 1:500 dilution of anti-polycystin-1 was used.

15 6.1.4. ADHESION ASSAYS

Epithelial cell adhesion to extracellular matrix substrates was determined
 by plating of 1,000-4,000 cells per well of 96-well cluster plates precoated with matrix
 protein as indicated. Cells were allowed to adhere under serum-free conditions for
 varied lengths of time (4 hr.-48 hr.), non-adherent cells removed by washing and adherent
 20 cell number determined by a linear colorimetric assay (Promega Cell titer 96™AQ,

4-24 hr. incubation with 3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxy methoxyphenyl)-2-(4-sulphonyl)-2M-tetrazolium, (MTS) and phenazine methosulfate (PMS).

6.1.5 DOUBLE IMMUNOFLUORESCENT LABELING

5 Cells on coverslips were washed and fixed in freshly prepared, ice-cold 4% paraformaldehyde for 3 min, washed 3 times, 3 min each with PBS and then subjected to double-labeling with polyclonal anti-polycystin coupled to anti-rabbit-IgG-rhodamine and monoclonal anti- α 2 β 1-integrin (1:100, Chemicon), or vinculin (1:500, Sigma), or paxillin (Sigma, 1:250), coupled to anti- mouse-IgG-FITC.

10 6.1.6. WESTERN IMMUNOBLOT ANALYSIS

SDS, NP-40 and Triton X-100 lysates were prepared from subconfluent and confluent monolayers of cell cultures washed in PBS containing a comprehensive protease inhibitor cocktail and vanadate. Equal amounts of protein (20 mg) were loaded per lane and separated by SDS-PAGE, transferred to nitrocellulose or nylon membranes and blotted with anti-polycystin (1:12,500), anti-phosphotyrosine, RC-20, (Transduction Laboratories, 1:2,500), anti- α 2 integrin (Chemicon, 1:2,000); anti-actin (Chemicon, 1:2,500); anti-vinculin (Sigma, 1:7,500 ; anti-paxillin (Transduction, Laboratories, 1:10,000) and anti-FAK (Transduction Laboratories, 1:10,000).

6.1.7. IMMUNOPRECIPITATIONS

Immunoprecipitations were carried out on NP40/ Triton X-100 cell extracts (5 X 100 mg) containing protease inhibitors, using anti-PKD-1 and anti-phosphotyrosine antibodies added to precleared Protein A/G PLUS agarose beads for 1 hr. and 4°C. After 3 washes with immunoprecipitation buffer the pellet was resuspended in sample buffer and subjected to SDS-PAGE.

6.1.8 STATISTICS

Data were expressed as mean \pm standard error and groups were subjected to analysis of variance.

6.2 RESULTS

A 31 amino acid peptide polyclonal antibody, corresponding to amino acid residues 4161-4191 in polycystin-1 was raised in rabbits, affinity purified and characterized for its specificity (Figures 1 and 2). The anti-polycystin-1 antibody recognized a fusion protein containing the C-terminal, intracellular portion of polycystin-1 (residues 4105-4303) while the pre-immune serum did not (Figure 1A). In addition, a single band of anticipated >440kDa molecular weight was detected in cell lysates from human fetal collecting tubule cell lines, while pre-immune serum detected no bands (Figure 1B). This antibody was high titer both for Western immunoblot analysis where it was used at 1:12,500 dilution and also for immunohistochemical analyses where it was used at 1:500 on paraformaldehyde fixed, paraffin embedded tissue

sections (Figures 2A-C) and 1:1,000 on cultured cells (Figure 2D). Immunoreactivity was dramatically reduced or absent after competition by preadsorption with the purified peptide (Figures 2 E-H). No immunoreactivity was seen in tissue sections or cells after incubation in preimmune sera (Figures 3 D and H). This antibody could also be used for immunoprecipitation analysis detecting a >440kDa band which was absent from immunoprecipitations carried out with preimmune sera (Figure 10A).

Primary cell strains and immortalized cell lines of normal and ADPKD renal epithelia were characterized with regard to their polycystin-1 expression by immunocytochemical and Western immunoblot analyses (Figures 3 and 4). The cells selected for study also showed cell-type specific marker expression. Stationary, confluent cultures of fetal collecting tubule epithelia showed characteristically high levels of normal polycystin-1 (Figure 3E); normal adult collecting tubules showed low levels of expression (Figure 3F); and ADPKD epithelia showed high levels of expression of polycystin-1 (Figure 3G). ADPKD epithelia from 10 different kindreds showed high levels of expression by immunocytochemistry and Western immunoblot analysis, using our C-terminal region antibody. This was also the case in multiple independent immortalized clonal ADPKD cell lines established from a single sample known to contain a mutation at exon 16 and to predict a truncated protein (Harris, P.C. personal communication). This finding of high level expression of polycystin-1 protein expression in ADPKD cystic epithelia is consistent with the findings of several other laboratories using monoclonal and polyclonal antibodies raised against a variety of – terminal to C-terminal epitopes in human disease as well as in mice with a heterozygous PKD1

mutation introduced by targeted gene disruption (Ward et al., 1996, *Proc Natl Acad Sci USA* **93**:1524-1528; Griffin et al., 1996, *Proc Assoc Am Physicians* **108**:185-197; Peters et al., 1996, *Lab Invest* **75**:221-230; Geng et al., 1996, *J Clin Invest* **98**:2674-2682; Wilson, 1997, *Am J Physiol* **272**:434-F442; Van Adelsberg et al., 1997, *Am J Physiol* **272**:F451-F459; Weston et al., 1997, *Histochem J* **29**:847-856; Ong et al., 1999, *Am J Pathol* **154**:1721-1729; Lu et al., 1999, *Nat Genet* **21**:160-161). This expression pattern of polycystin-1, therefore, mirrored those seen in normal fetal, adult and ADPKD tissues (Figure 4A).

6.2.1 MATRIX ADHESION IN ADPKD

Previous studies have identified a variety of structural abnormalities in the extracellular matrix of human ADPKD kidneys and in matrix protein synthesis and turnover of cultured ADPKD epithelial cells *in vitro* (Wilson and Falkenstein, 1995, The Pathology of human renal cystic disease. In: Dodd SM, editor. Current topics in pathology, vol **88**. Heidelberg: Springer-Verlag, 1-50; Wilson et al., 1996, Pathogenesis of polycystic kidney disease: Altered cellular function. In: Watson ML and Torres VE, editors. Polycystic kidney disease. Oxford: Oxford Medical Publications ; Liu et al., 1992, *Am J Physiol* **263**:F697-F704). In initial studies, it was also noted that ADPKD epithelia were more adherent *in vitro* than their normal counterparts and that the degree of adhesion influenced subsequent epithelial cell proliferation (Wilson and Sherwood, 1991, *Kidney Int* **39**:450-463; Wilson et al., 1992, *J Cell Physiol* **150**:360-369). These observations have now been re-examined, expanded and evaluated with respect to

cell-type and matrix protein specificity using a highly reproducible, quantitative adhesion assay (Figure 5). Figure 5A is a typical representative profile of adhesion for primary epithelial cell strains derived from different segments of the nephron. Twenty separate experiments were conducted at 12-24 and 48 hour time-points, plating 2,000 and 1,000 cells respectively. An additional set of 12 experiments was carried out measuring adhesion after 15 min, 30 min, 90 min and 4 hr. adhesion plating 2,000-5,000 cells per well. It was consistently observed that ADPKD cell lines expressing high levels of polycystin-1 (Figure 5A, lane 6) were significantly (2 to 3-fold) more adherent than fetal or adult proximal tubule epithelia (PT Figure 5A, lanes 1 and 3), adult thick ascending limb epithelia (TAL, Figure 5A, lane 4) or adult collecting tubule epithelia (CT, Figure 5A, lane 5) expressing lower levels of normal polycystin-1. Fetal collecting tubule epithelia (HFCT) expressed intermediate to high levels of polycystin-1 (Figure 3E, Figure 4B, lanes 1 and 2) and showed intermediate levels of adhesion to type I collagen (Figure 5A, lane 2).

The specificity of this adhesion preference was demonstrated by a differential effect with regard to the type of extracellular matrix protein used in the assay. Adherence to uncoated tissue culture plastic ("none") and to type I collagen were the most discriminatory and significantly greater than adhesion to type IV collagen or to fibronectin (Figure 5B). No difference in the adhesion of normal and ADPKD cells was seen on laminin substrates. This finding was of interest since type I collagen is a major component of the immediate extracellular matrix of both ADPKD and fetal ureteric bud epithelia, *in vivo* (Ekblom, 1989, *FASEB J* 3:2141-2150). By contrast, normal adult renal

epithelia are in immediate contact with an intact basement membrane, which contains type IV collagen and laminin, not type I collagen which is restricted to the interstitium. This 2 to 3-fold increase in adhesion to type I collagen by ADPKD epithelia was also seen in conditionally immortalized ADPKD clonal cell lines (Figure 5C, lanes 6, 7 and 8), suggesting this is a widespread cellular alteration in ADPKD cyst lining epithelia. Again, HFCT conditionally immortalized clones showed higher levels of attachment than other normal renal epithelial cell types (Figure 5C, lane 3). These results show that the same patterns of differential adhesion to type 1 collagen were retained in conditionally immortalized cell lines derived from human renal epithelia cultured at the non-permissive temperature of 37°C, conditions under which normal differentiation characteristics have been demonstrated by marker analysis (Racusen, 1995, *Kidney Int* 48:536-543; Wilson 1999, *In vitro* methods in renal research. In: Barrat TM, et al., editors. Pediatric nephrology, 4th ed. Baltimore: Lippincott Williams & Wilkins, 269-281).

6.2.2. ROLE FOR $\alpha 2\beta 1$ INTEGRIN IN ADHESION

A potential role for $\alpha 2\beta 1$ integrin in this adhesion difference was studied since this heterodimeric complex has been shown to act as a receptor for type I collagen (Hynes, 1992, *Cell* 69:11-25). Immunocytochemical analysis showed that ADPKD epithelia grown to confluence on type I collagen, stained intensely for $\alpha 2\beta 1$ integrin and demonstrated a focal pattern of distribution characteristic of accumulation at points of cell contact with its substratum, seen as ridges of reaction product at the plane of cell-substrate contact (Figure 6A). In addition, an approximately 2-fold increase in

a2-integrin content in adherent ADPKD versus adherent normal epithelia was seen by Western immunoblot analysis of whole cell extracts (Figure 6B, lanes 8 and 9) potentially implicating an integrin-mediated mechanism for the increased adhesive properties of ADPKD epithelia. Functional involvement of $\alpha 2\beta 1$ integrin in polycystin-related adhesion was also suggested by inhibition of adhesion of ADPKD epithelial cells by a blocking antibody against $\alpha 2\beta 1$ integrin, "6F1"(Coller et al, 1989, Blood 74:182-192; Figure 6C lanes 8, 9 and 10) but not of normal cells (Figure 6C, lines 2, 4 and 6).

Western immunoblot analysis of intracellular proteins which are known to interact with $\alpha 2\beta 1$ integrin in other cell systems, also showed some trends to quantitative changes in total cell expression levels in confluent cells when levels in normal epithelia were compared with those in ADPKD epithelia after plating of the same numbers of cells on type I collagen and allowing adhesion for identical lengths of time. Vinculin (Figure 7A) and paxillin (Figure 7B) were approximately 2-fold increased in ADPKD cells (lanes 4 and 5), while focal adhesion kinase (pp125^{FAK}) was apparently decreased in ADPKD epithelia by comparison to normal renal epithelial cells (Figure 7C, lanes 4 and 5).

6.2.3 COLOCALIZATION OF POLYCYSTIN-1 WITH FOCAL ADHESION PLAQUE PROTEINS

The finding of a specific alteration of $\alpha 2\beta 1$ -integrin function and quantitative increases in $\alpha 2\beta 1$ -integrin, vinculin and paxillin in ADPKD epithelial cells suggested the possibility that polycystin-1 might interact directly with $\alpha 2\beta 1$ -integrin at

sites of cell attachment to the underlying matrix. To test this notion, double labeling immunofluorescence and confocal microscopy was carried out using anti-polycystin-1 antibody raised in rabbit and anti- $\alpha 2\beta 1$ -integrin mouse monoclonal antibody. These studies (Figure 8A-C) using rhodamine conjugated secondary antibody for polycystin-1 and fluorescein isothiocyanate (FITC) conjugated secondary antibody for $\alpha 2\beta 1$ -integrin, demonstrated complete co-localization of these proteins in focal regions of the basal cell surface in the plane of attachment with the substratum.

Adhesion of cells to a matrix *in vitro* involves the transient formation of focal clusters, to which structural and signal transduction proteins are recruited, including integrins, vinculin, paxillin, and pp^{125FAK} (Clark and Brugge, 1995, *Science* **268**:233-239). In fibroblasts, classical, mature focal adhesions are typically needle-shaped in appearance. In epithelial cells, however, larger, more globular structures have reported to form, particularly in the initial stages of adhesion to a substratum (Jockusch et al., 1995, *Rev Cell Dev Biol* **11**:379-416). Double labeling immunofluorescence studies showed completely overlapping co-localization of polycystin-1 with vinculin (Figure 8D-F) and paxillin (Figure 8G-I) which were detected together with $\alpha 2\beta 1$ -integrin in large focal clusters at the cell membrane at points of contact with the matrix. Similarly, co-localizations of polycystin-1, $\alpha 2\beta 1$ -integrin, vinculin and paxillin were demonstrated in focal clusters in cultures of normal human fetal collecting tubules after short periods of adhesion to type I collagen (Figure 9). In summary, these co-localization studies establish identical distributions of PKD-1 with $\alpha 2\beta 1$ -integrin, vinculin and paxillin within cluster-like structures at the cell-matrix interface in response to initial adherence to

type I collagen matrix apparently both in normal and ADPKD epithelia. It should be noted, however, that, consistent with the findings presented in Figure 5, many more ADPKD epithelia were adherent and showed cluster formation after 4 hours than fetal collecting tubule epithelia.

5 To establish whether there was direct physical association between PKD-1 and these co-distributing proteins, cell extracts were immunoprecipitated with anti-polycystin-1 antibodies and associated proteins examined by immunodetection using Western blotting (Figure 10). In normal fetal collecting tubule epithelia (Figure 10B), co-immunoprecipitation of PKD-1 with vinculin (lane 2), paxillin (lane 3) and pp125^{FAK} (lane 4) was demonstrated. Interestingly, in ADPKD epithelia (Figure 10C), although
10 vinculin (lane 2) and paxillin (lane 3) were present in the co-immunoprecipitates with PKD-1, pp125^{FAK} (lane 4) was not apparently associated with polycystin-1 protein complexes. This therefore suggested an alteration in tyrosine phosphorylation of polycystin-1 might play a role in alterations of adhesion in ADPKD epithelia

15 6.2.4 TYROSINE PHOSPHORYLATION OF POLYCYSTIN-1

 The deduced amino acid sequence of polycystin-1 suggests a potential site for tyrosine phosphorylation at Tyr₄₁₂₇ in the cytoplasmic C-terminal domain of the protein (International PKD Consortium, 1995, *Cell* **81**:289-298). To determine biochemically, whether polycystin-1 is indeed a target of tyrosine phosphorylation, two
20 types of experiment were carried out. In the first set, parallel samples of cells were

subjected to immunoblot analyses using anti-phosphotyrosine and anti-polycystin-1 (C-terminal domain) antibodies (Figures 11A, B). Using actively proliferating, adherent human renal epithelial primary and immortalized cell lines with detectable levels of polycystin-1 by Western blot analysis (Figure 9A), it was shown by parallel analysis of anti-phosphotyrosine immunoreactivity on stripped blots, that polycystin-1 was tyrosine phosphorylated (Figure 11B). As predicted, additional proteins in addition to polycystin-1 were also tyrosine phosphorylated as represented by lower molecular weight bands. The same results were obtained if anti-PKD1 blots were stripped and reprobed with anti-phosphotyrosine antibodies.

Definitive evidence that polycystin-1 can be phosphorylated on tyrosine residues was obtained by immunoprecipitation of ADPKD and normal renal epithelial cell proteins with anti-phosphotyrosine antibodies followed by immunoblot analysis with anti-polycystin-1 antibody. A major band of > 440kDa was detected in proliferative and adherent normal and ADPKD cells, indicating the tyrosine phosphorylation of polycystin-1 in these cells (Figure 11C). Recently we have extended these findings by demonstration of tyrosine phosphorylation of the PKD-1- C-terminal domain fusion protein *in vitro* (Li et al., 1999, *Am J Physiol* **263**:F697-F704). Of interest, the level of tyrosine phosphorylation in the ADPKD epithelial cells appeared to be lower than in normal cells (Figure 11C lane 1) suggesting that reduction in polycystin-1 tyrosine phosphorylation may be an important consequence of *PKD1* gene mutation(s). This is consistent with the majority of mutations reported to date being deletions; missense or introduction of premature stop codons which would result in a truncated protein and

removal of several putative tyrosine phosphorylation sites. It has recently shown that the phosphorylation of the PKD1-C-terminal domain fusion protein *in vitro* by c-src can be abrogated entirely by site directed mutation at tyrosine position Y 4237 (Li et al., 1999, *Am J Physiol* **263**:F697-F704).

5 Various publications are cited herein which are hereby incorporated, by reference, in their entirety.

 The present invention is not to be limited in scope by the embodiments disclosed in the examples which are intended as an illustration of one aspect of the invention, and any compositions or methods which are functionally equivalent are within
10 the scope of this invention. Indeed, various modifications of the invention in addition to those shown and described herein will become apparent to those skilled in the art from the foregoing description. Such modifications are intended to fall within the scope of the claims.

What is claimed:

1. A method for identifying a compound capable of modulating polycystin-1 activity, comprising:
 - (a) contacting a test compound to a recombinantly engineered cell expressing a polycystin-1 protein, or a variant thereof, resulting in expression of a mutant cell phenotype;
 - (b) measuring the expression of the mutant phenotype; and
 - (c) comparing the level of expression of the mutant phenotype obtained in (b) to the level of expression of a mutant phenotype obtained in the presence of a vehicle control: such that if the level obtained in (b) differs from that obtained in the presence of a vehicle control, a compound capable of modulating polycystin-1 activity has been identified.
2. The method of Claim 1 wherein the mutant phenotype is an increase in cell adherence to type I collagen coated substrates.
3. The method of Claim 1 wherein the mutant phenotype is an increase in apical expression of NaK-ATPase on the cell membrane.
4. The method of Claim 1 wherein the mutant phenotype is increased expression of β -2-NaK-ATPase within the cell.

5. The method of Claim 4 wherein the expression of β -2-NaK-ATPase within the cell is measured using an anti- β -2-NaK-ATPase antibody.
6. The method of Claim 1 wherein the mutant phenotype is decreased incorporation of focal adhesion kinase into focal adhesion complexes.
- 5 7. The method of Claim 6 wherein the incorporation of focal adhesion kinase into focal adhesion complexes is measured using an anti-focal adhesion kinase antibody.
8. The method of Claim 1 wherein the recombinantly engineered cell further comprises an epitope tagged polycystin-1 interacting protein.
- 10 9. The method of Claim 1 wherein the recombinantly engineered cell further comprises an epitope tagged focal adhesion kinase protein.
10. The method of Claim 1 wherein the polycystin-1 protein, or variant thereof, is epitope tagged.
11. A method for identifying a compound capable of modulating polycystin-2 activity, comprising:
- 15

- 5 (a) contacting a test compound to a recombinantly engineered cell expressing a polycystin-2 protein, or variant thereof, resulting in expression of a mutant cell phenotype;
- (b) measuring the expression of the mutant phenotype; and
- 10 (c) comparing the level of expression of mutant phenotype obtained in (b) to the level of expression of a mutant phenotype obtained in the presence of a vehicle control: such that if the level obtained in (b) differs from that obtained in the presence of a vehicle control, a compound capable of modulating polycystin-2 activity has been identified.

12. The method of Claim 11 wherein the mutant phenotype is an increase in cell adherence to type I collagen coated substrates.

13. The method of Claim 12 wherein the mutant phenotype is an increase in apical expression of NaK-ATPase on the cell membrane.

15 14. The method of Claim 13 wherein the mutant phenotype is increased expression of β -2-NaK-ATPase within the cell.

15. The method of Claim 14 wherein the expression of β -2-NaK-ATPase within the cell is measured using an anti- β -2-NaK-ATPase antibody.

16. The method of Claim 11 wherein the mutant phenotype is decreased incorporation of focal adhesion kinase into focal adhesion complexes.
17. The method of Claim 16 wherein the incorporation of focal adhesion kinase into focal adhesion complexes is measured using an anti-focal adhesion kinase antibody.
18. The method of Claim 11 wherein the recombinantly engineered cell further comprises an epitope tagged polycystin-2 interacting protein.
19. The method of Claim 11 wherein the recombinantly engineered cell further comprises an epitope tagged focal adhesion kinase protein.
20. The method of Claim 11 wherein the polycystin-2 protein, or variant thereof, is epitope tagged.

ABSTRACT OF THE INVENTION

The present invention provides cell-based screening assays designed to identify agents that regulate the activity of the polycystic kidney disease proteins encoded by the PKD-1 and PKD-2 genes and that may be useful in the treatment of polycystic kidney disease. The assays of the invention comprise the contacting of genetically engineered cells expressing a mutant or truncated PKD gene product with a test agent and assaying for a decrease in the PKD mediated mutant phenotype. Characteristics associated with such a mutant phenotype include increased adherence to type I collagen coated surfaces; apical expression of NaK-ATPase on the cell membrane; increased expression of β -2-NaK-ATPase; and decreased focal adhesion kinase (FAK) incorporation into focal adhesion complexes, and inability to form tubular structures in a gel matrix. To facilitate the screening methods of the invention, cells may be genetically engineered to express epitope tagged PKD gene products and/or epitope tagged PKD interacting proteins (PKD-IP). Such interacting proteins include, for example, focal adhesion complex proteins such as FAK, paxillin, vinculin, talin and the like.

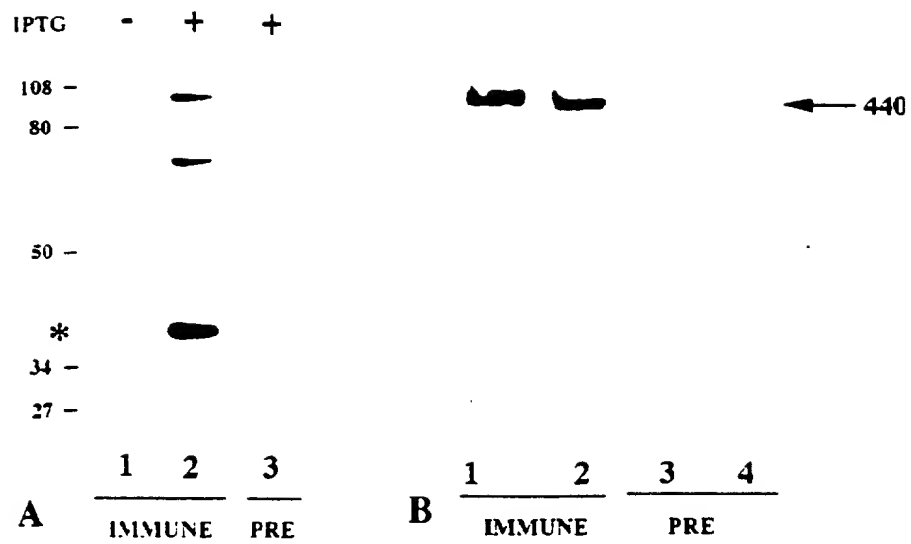


FIGURE 1A-B

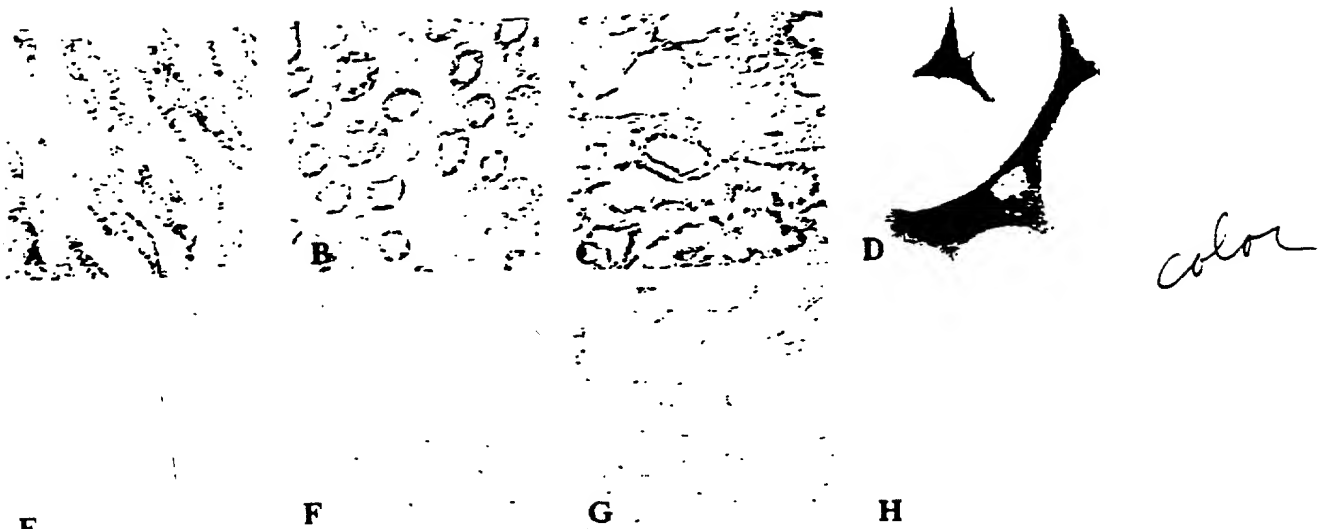


FIGURE 2A-H

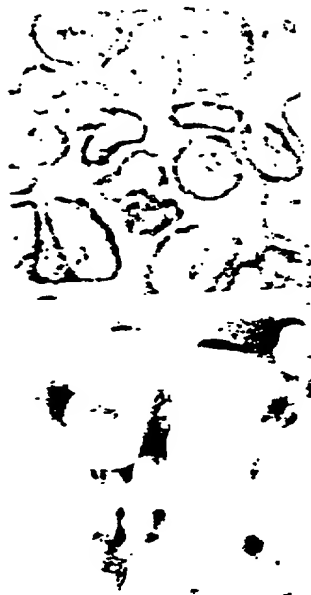
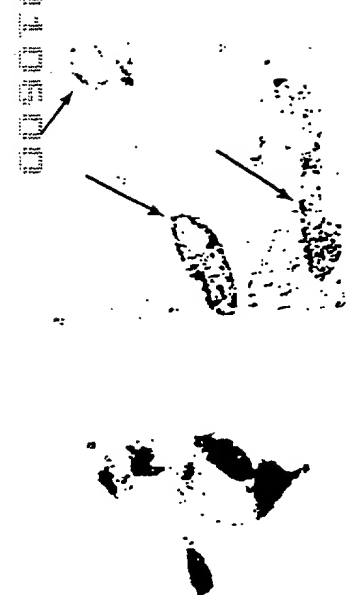
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E

F

G

H



color

FIGURE 3A-H

Wilson et al

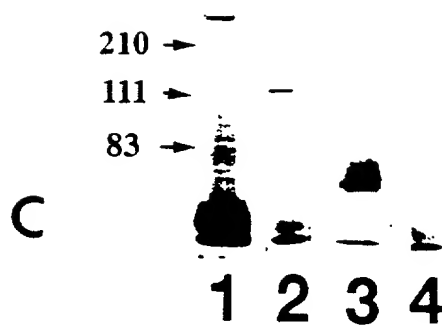
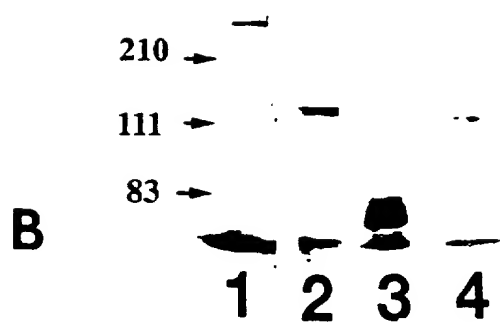
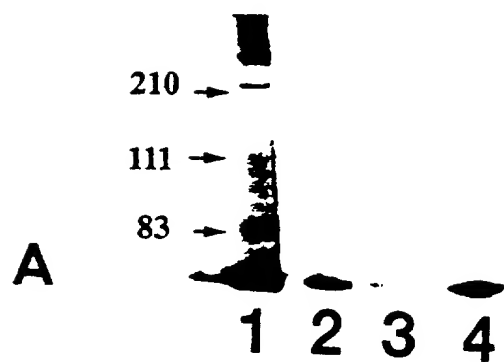


FIGURE 4A-C

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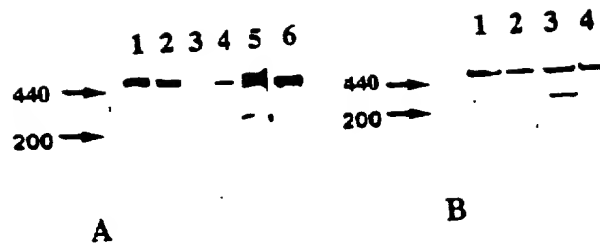


FIGURE 5A-B

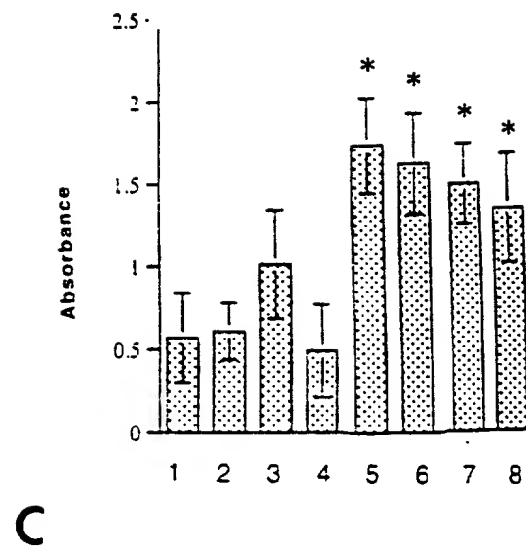
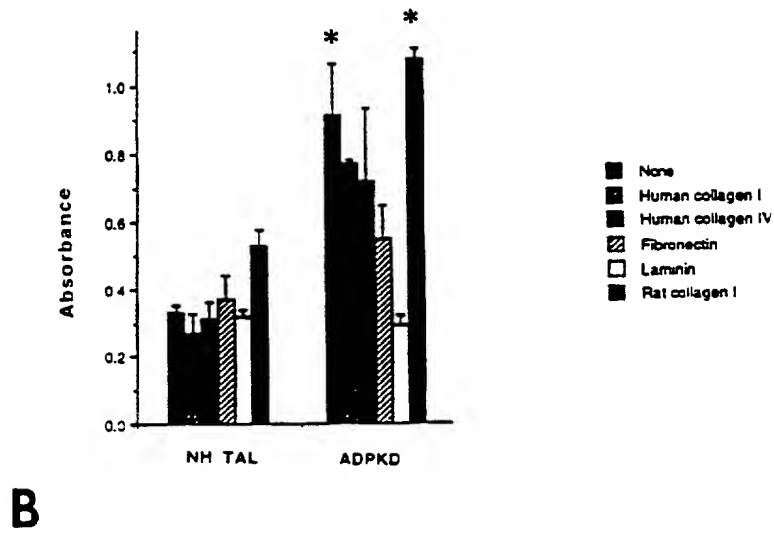
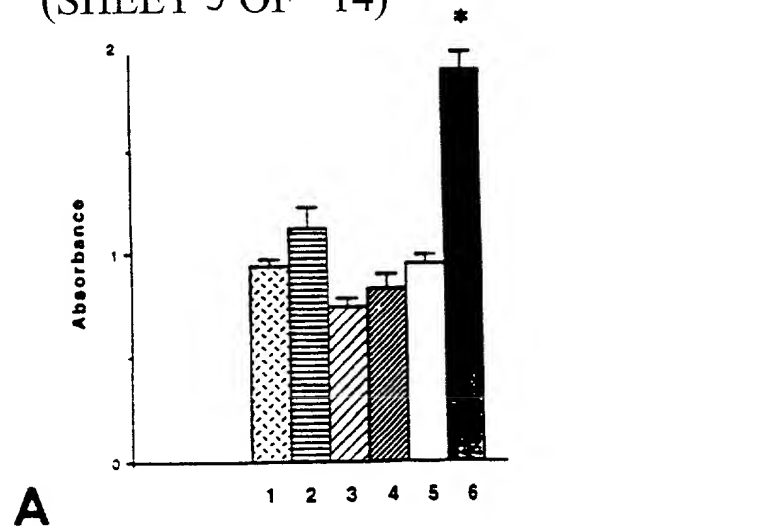


FIGURE 6A-C

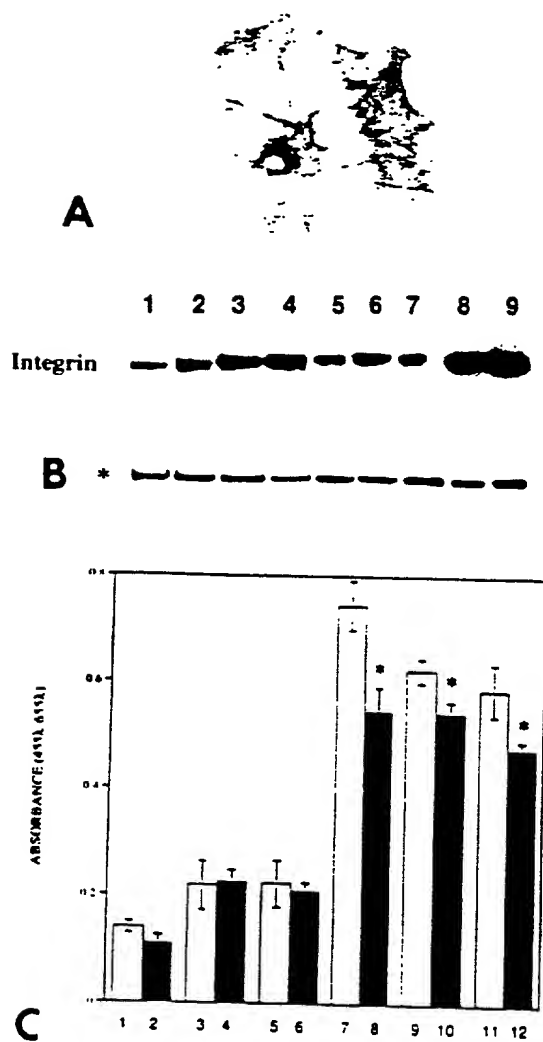
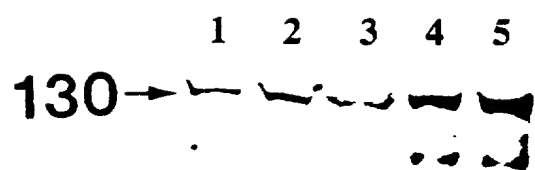
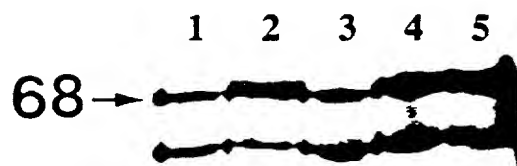


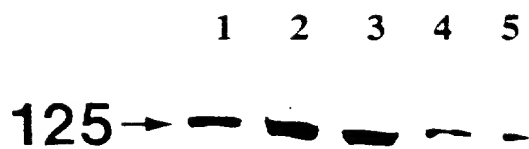
FIGURE 7A-C



A



B

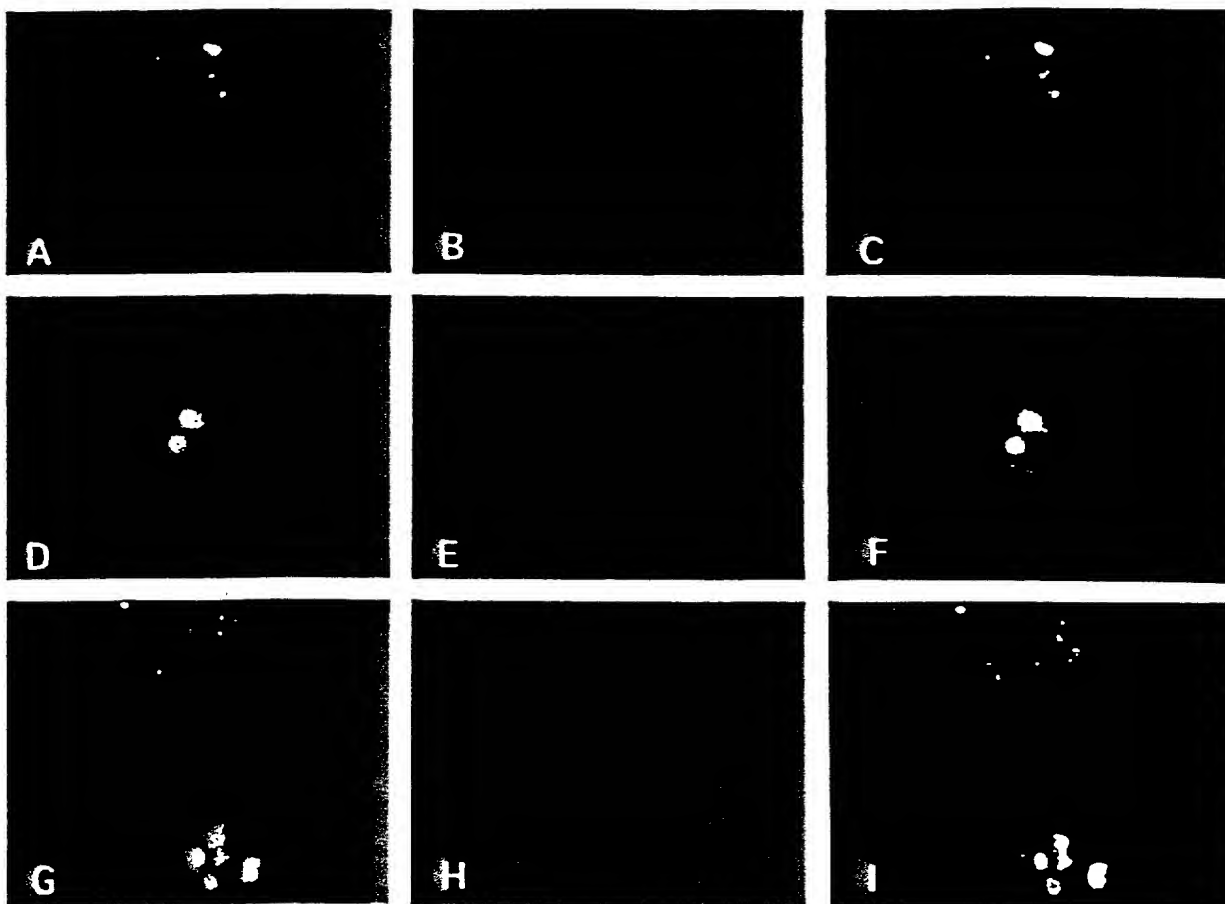


C

FIGURE 8A-C

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W1000-10-11



Color

FIGURE 9A-I

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PKD1 Protein in Epithelial Attachment

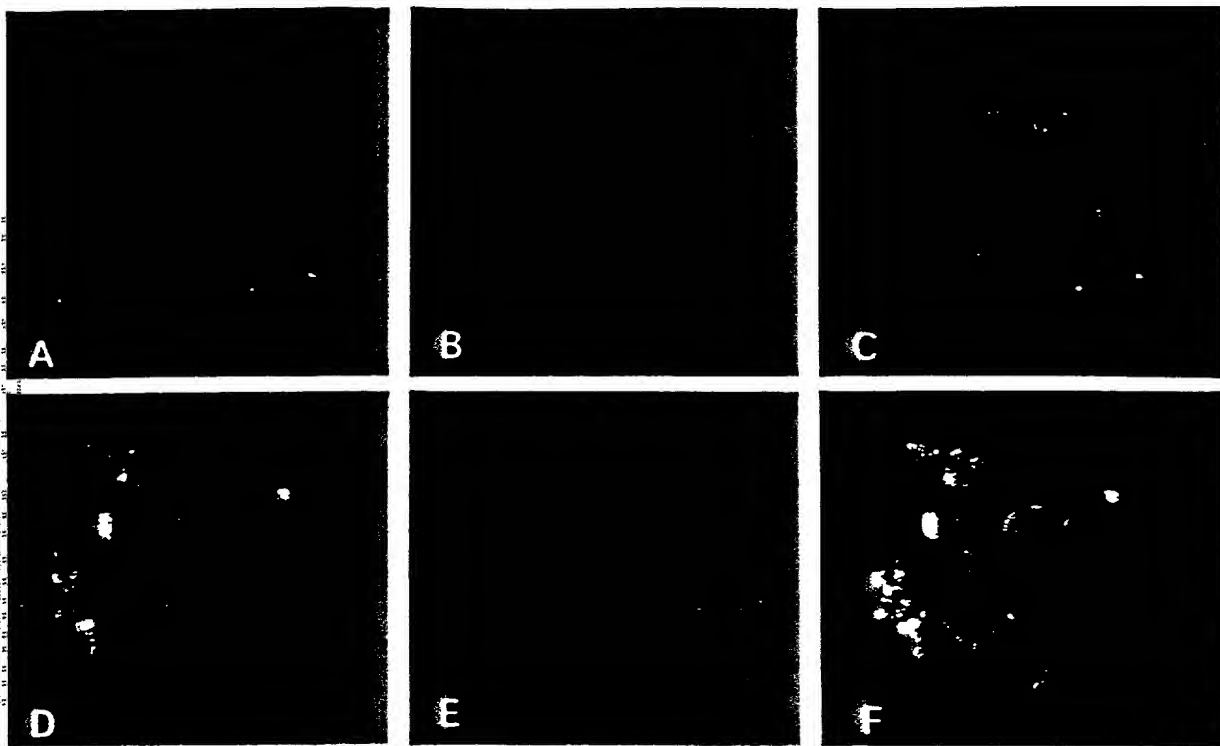


FIGURE 10A-F

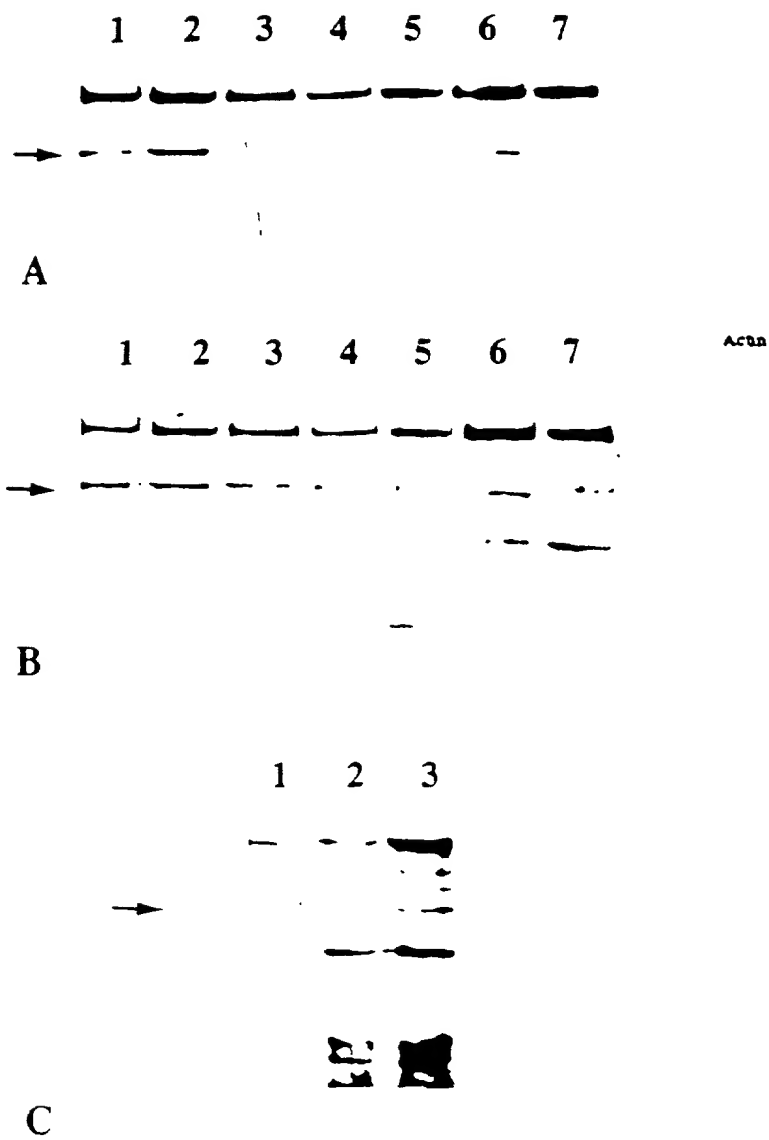
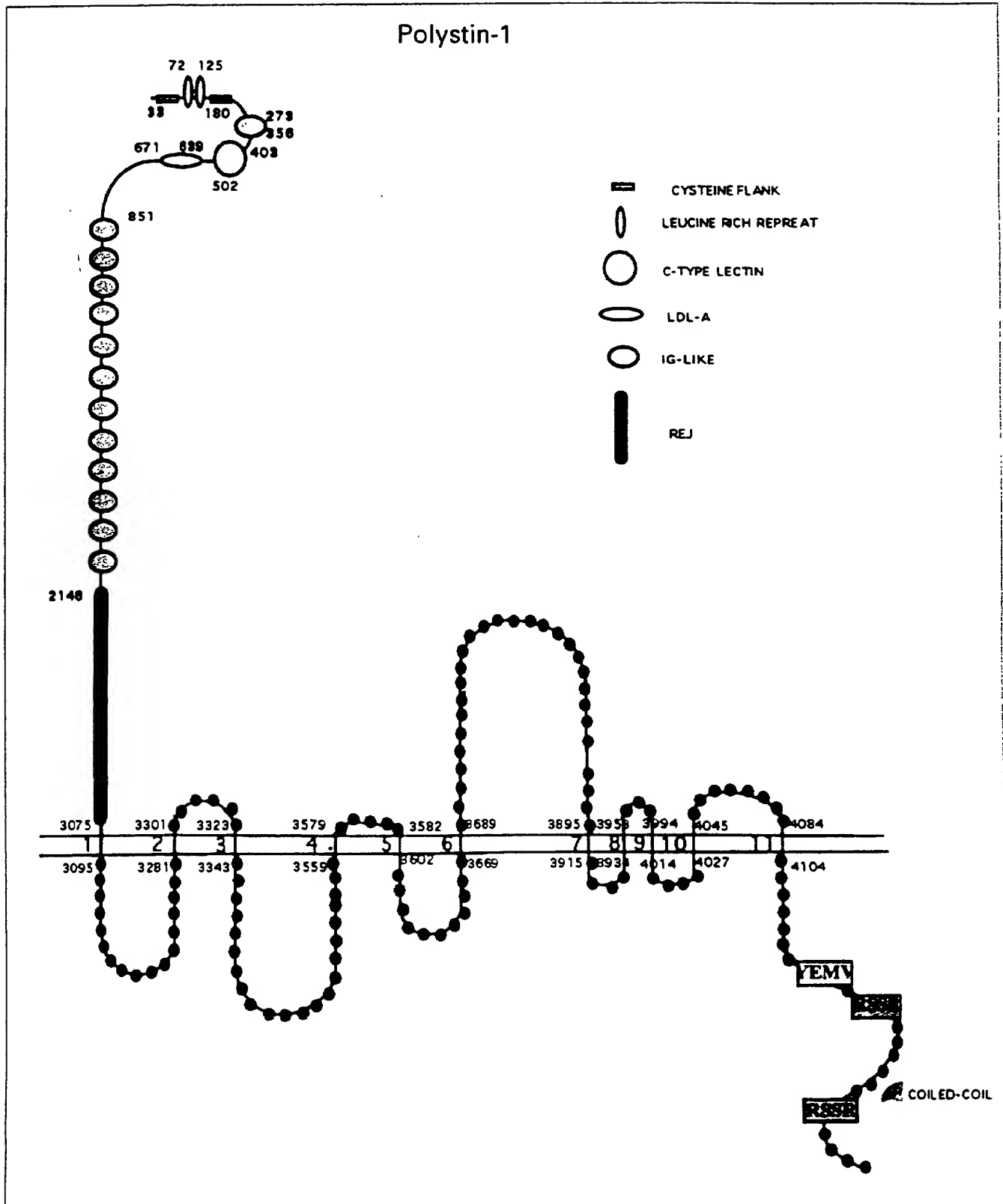


FIGURE 11A-C

FIGURE 12 A



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FIGURE 12 B

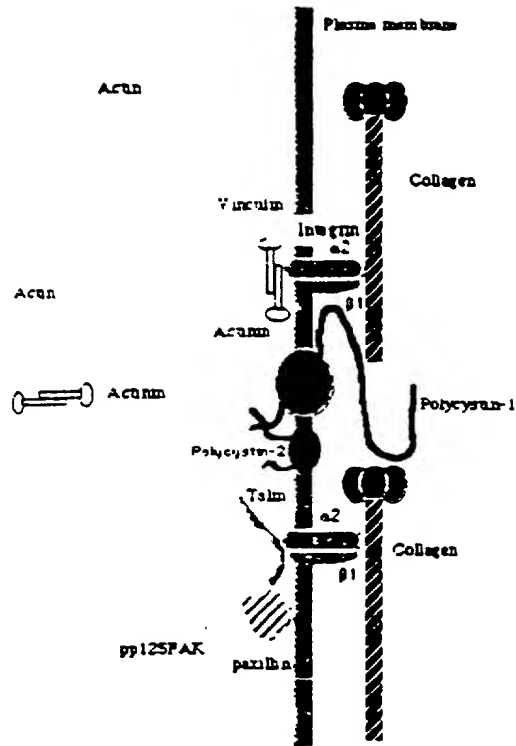
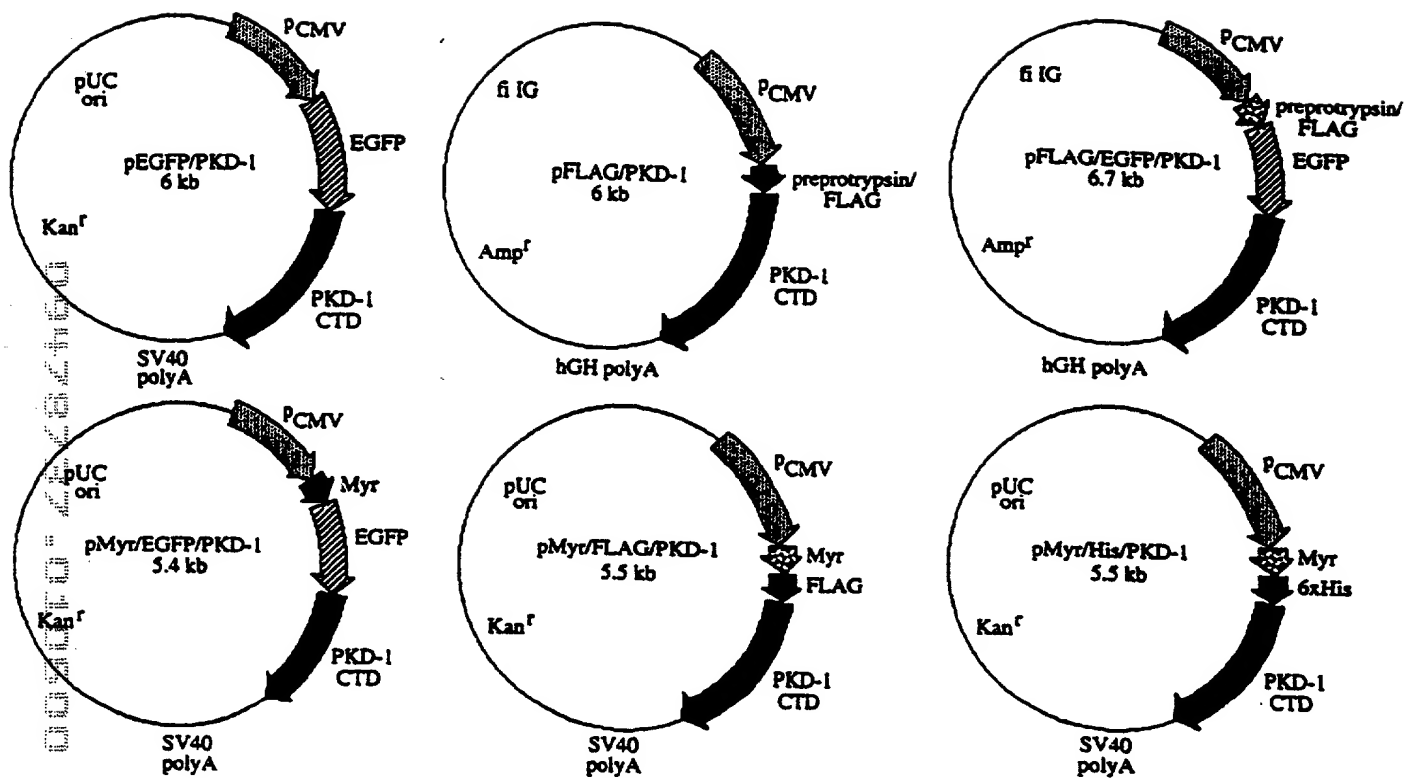


FIGURE 12 C

FIGURE 13



**COMBINED DECLARATION
AND POWER OF ATTORNEY**

(Original, Design, National Stage of PCT, Divisional, Continuation or C-I-P Application)

As a below named inventor, I hereby declare that:

My residence, post office address and citizenship are as stated below next to my name; I believe I am the original, first and sole inventor (if only one name is listed below) or an original, first and joint inventor (if plural names are listed below) of the subject matter which is claimed and for which a patent is sought on the invention entitled:

**"SCREENING METHODS FOR COMPOUNDS USEFUL IN THE
REGULATION OF POLYCYSTIC KIDNEY DISEASE"**

This declaration is of the following type:

- ☒ original
☐ design
☐ national stage of PCT.
☐ divisional
☐ continuation
☐ continuation-in-part (C-I-P)

the specification of which: *(complete (a), (b), or (c))*

- (a) ☒ is attached hereto.
(b) ☐ was filed on as Application Serial No. and was amended on *(if applicable)*.
(c) ☐ was described and claimed in PCT International Application No. filed on and was amended on *(if applicable)*.

Acknowledgement of Review of Papers and Duty of Candor

I hereby state that I have reviewed and understand the contents of the above identified specification, including the claims, as amended by any amendment referred to above.

I acknowledge the duty to disclose information which is material to the patentability of the subject matter claimed in this application in accordance with Title 37, Code of Federal Regulations § 1.56.

☐ In compliance with this duty there is attached an information disclosure statement. 37 CFR 1.98.

Priority Claim

I hereby claim foreign priority benefits under Title 35, United States Code, § 119(a)-(d) of any foreign application(s) for patent or inventor's certificate or of any PCT International Application(s) designating at least one country other than the United States of America listed below and have also identified below any foreign application(s) for patent or inventor's certificate or any PCT International Application(s) designating at least one country other than the United States of America filed by me on the same subject matter having a filing date before that of the application on which priority is claimed

(complete (d) or (e))

- (d) ☒ no such applications have been filed.
(e) ☐ such applications have been filed as follows:

PRIOR FOREIGN/PCT APPLICATION(S) FILED WITHIN 12 MONTHS (6 MONTHS FOR DESIGN) PRIOR TO SAID APPLICATION			
COUNTRY	APPLICATION NO.	DATE OF FILING (day, month, year)	DATE OF ISSUE (day, month, year)
			PRIORITY CLAIMED UNDER 35 USC 119 <input type="checkbox"/> YES <input type="checkbox"/> NO <input type="checkbox"/>
			<input type="checkbox"/> YES <input type="checkbox"/> NO <input type="checkbox"/>
			<input type="checkbox"/> YES <input type="checkbox"/> NO <input type="checkbox"/>
ALL FOREIGN APPLICATION(S), IF ANY, FILED MORE THAN 12 MONTHS (6 MONTHS FOR DESIGN) PRIOR TO SAID APPLICATION			
			<input type="checkbox"/> YES <input type="checkbox"/> NO <input type="checkbox"/>
			<input type="checkbox"/> YES <input type="checkbox"/> NO <input type="checkbox"/>
			<input type="checkbox"/> YES <input type="checkbox"/> NO <input type="checkbox"/>

Claim for Benefit of Prior U.S. Provisional Application(s)

I hereby claim the benefit under Title 35, United States Code, § 119(e) of any United States provisional application(s) listed below:

Provisional Application Number	Filing Date

Claim for Benefit of Earlier U.S./PCT Application(s) under 35 U.S.C. 120

(complete this part only if this is a divisional, continuation or C-I-P application)

I hereby claim the benefit under Title 35, United States Code, § 120 of any United States application(s) or PCT international application(s) designating the United States of America that is/are listed below and, insofar as the subject matter of each of the claims of this application is not disclosed in the prior application(s) in the manner provided by the first paragraph of Title 35, United States Code § 112, I acknowledge the duty to disclose information as defined in Title 37, Code of Federal Regulations, § 1.56 which occurred between the filing date of the prior application(s) and the national or PCT international filing date of this application:

Application Serial No.)	(Filing Date)	(Status) (patented, pending, abandoned)
Application Serial No.)	(Filing Date)	(Status) (patented, pending, abandoned)

Power of Attorney

As a named inventor, I hereby appoint Dana M. Raymond, Reg. No. 18,540; Frederick C. Carver, Reg. No. 17,021; Francis J. Hone, Reg. No. 18,662; Joseph D. Garon, Reg. No. 20,420; Arthur S. Tenser, Reg. No. 18,839; Ronald B. Hildreth, Reg. No. 19,498; Thomas R. Nesbitt, Jr., Reg. No. 22,075; Robert Neuner, Reg. No. 24,316; Richard G. Berkley, Reg. No. 25,465; Richard S. Clark, Reg. No. 26,154; Bradley B. Geist, Reg. No. 27,551; James J. Maune, Reg. No. 26,946; John D. Murnane, Reg. No. 29,836; Henry Tang, Reg. No. 29,705; Robert C. Scheinfeld, Reg. No. 31,300; John A. Fogarty, Jr., Reg. No. 22,348; Louis S. Sorell, Reg. No. 32,439; Rochelle K. Seide Reg. No. 32,300; Gary M. Butter, Reg. No. 33,841; Marta E. Delsignore, Reg. No. 32,689; and Lisa B. Kole, Reg. No. 35,225 of the firm of BAKER & BOTTS, L.L.P., with offices at 30 Rockefeller Plaza, New York, New York 10112, as attorneys to prosecute this application and to transact all business in the Patent and Trademark Office connected therewith

SEND CORRESPONDENCE TO: BAKER & BOTTS, L.L.P. 30 ROCKEFELLER PLAZA, NEW YORK, N.Y. 10112 CUSTOMER NUMBER: 21003	DIRECT TELEPHONE CALLS TO: BAKER & BOTTS, L.L.P. (212) 705-5000
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I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section

1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

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POST OFFICE ADDRESS	POST OFFICE ADDRESS 525 East 80 th Street, Apt #8D	CITY New York	STATE or COUNTRY New York	ZIP CODE 10021
DATE x 12/22/99	SIGNATURE OF INVENTOR x <i>Christopher R. Burrow</i>			
FULL NAME OF THIRD JOINT INVENTOR, IF ANY	LAST NAME	FIRST NAME	MIDDLE NAME	
RESIDENCE & CITIZENSHIP	CITY	STATE or FOREIGN COUNTRY	COUNTRY OF CITIZENSHIP	
POST OFFICE ADDRESS	POST OFFICE ADDRESS	CITY	STATE or COUNTRY	ZIP CODE
DATE	SIGNATURE OF INVENTOR			
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RESIDENCE & CITIZENSHIP	CITY	STATE or FOREIGN COUNTRY	COUNTRY OF CITIZENSHIP	
POST OFFICE ADDRESS	POST OFFICE ADDRESS	CITY	STATE or COUNTRY	ZIP CODE
DATE	SIGNATURE OF INVENTOR			
FULL NAME OF FIFTH JOINT INVENTOR, IF ANY	LAST NAME	FIRST NAME	MIDDLE NAME	
RESIDENCE & CITIZENSHIP	CITY	STATE or FOREIGN COUNTRY	COUNTRY OF CITIZENSHIP	
POST OFFICE ADDRESS	POST OFFICE ADDRESS	CITY	STATE or COUNTRY	ZIP CODE
DATE	SIGNATURE OF INVENTOR			
FULL NAME OF SIXTH JOINT INVENTOR, IF ANY	LAST NAME	FIRST NAME	MIDDLE NAME	
RESIDENCE & CITIZENSHIP	CITY	STATE or FOREIGN COUNTRY	COUNTRY OF CITIZENSHIP	
POST OFFICE ADDRESS	POST OFFICE ADDRESS	CITY	STATE or COUNTRY	ZIP CODE
DATE	SIGNATURE OF INVENTOR			